

Advances in Anatomy, Embryology and Cell Biology

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# Sensing the Environment: Regulation of Local and Global Homeostasis by the Skin's Neuroendocrine System

Reviews and critical articles covering the entire field of normal anatomy (cytology, histology, cyto- and histochemistry, electron microscopy, macroscopy, experimental morphology and embryology and comparative anatomy) are published in *Advances in Anatomy, Embryology and Cell Biology*. Papers dealing with anthropology and clinical morphology that aim to encourage cooperation between anatomy and related disciplines will also be accepted. Papers are normally commissioned. Original papers and communications may be submitted and will be considered for publication provided they meet the requirements of a review article and thus fit into the scope of "Advances". English language is preferred.

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With 23 figures

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## Abstract

Skin, the body's largest organ, is strategically located at the interface with the external environment where it detects, integrates, and responds to a diverse range of stressors including solar radiation. It has already been established that the skin is an important peripheral neuro-endocrine-immune organ that is tightly networked to central regulatory systems. These capabilities contribute to the maintenance of peripheral homeostasis. Specifically, epidermal and dermal cells produce and respond to classical stress neurotransmitters, neuropeptides, and hormones. Such production is stimulated by ultraviolet radiation (UVR), biological factors (infectious and noninfectious), and other physical and chemical agents. Examples of local biologically active products are cytokines, biogenic amines (catecholamines, histamine, serotonin, and *N*-acetyl-serotonin), melatonin, acetylcholine, neuropeptides including pituitary (proopiomelanocortin-derived ACTH,  $\beta$ -endorphin or MSH peptides, thyroid-stimulating hormone) and hypothalamic (corticotropin-releasing factor and related urocortins, thyroid-releasing hormone) hormones as well as enkephalins and dynorphins, thyroid hormones, steroids (glucocorticoids, mineralocorticoids, sex hormones, 7- $\delta$  steroids), secosteroids, opioids, and endocannabinoids. The production of these molecules is hierarchical, organized along the algorithms of classical neuro-endocrine axes such as hypothalamic-pituitary-adrenal axis (HPA), hypothalamic-thyroid axis (HPT), serotonergic, melatonergic, catecholaminergic, cholinergic, steroid/secosteroidogenic, opioid, and endocannabinoid systems. Dysregulation of these axes or of communication between them may lead to skin and/or systemic diseases. These local neuroendocrine networks are also addressed at restricting maximally the effect of noxious environmental agents to preserve local and consequently global homeostasis. Moreover, the skin-derived factors/systems can also activate cutaneous nerve endings to alert the brain on changes in the epidermal or dermal environments, or alternatively to activate other coordinating centers by direct (spinal cord) neurotransmission without brain involvement. Furthermore, rapid and reciprocal communications between epidermal and dermal and adnexal compartments are also mediated by neurotransmission including antidromic modes of conduction. In conclusion, skin cells and skin as an organ coordinate and/or regulate not only peripheral but also global homeostasis.



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# List of Abbreviations

11 $\beta$ -HSD	11 $\beta$ -Hydroxysteroid dehydrogenase
2-AG	2-Arachidonoylglycerol
5-HT	5-Hydroxytryptamine, serotonin
5-HT <sub>1-7</sub>	5-HT receptors types 1–7
5-HTOL	5-Hydroxytryptophanol
5HTT	Serotonin transporter
5MTOL	5-Methoxytryptophol
5MTT	5-Methoxytryptamine
6BH4	6R-L-erythro-5, 6,7,8-tetrahydrobiopterin
7DHC	7-Dehydrocholesterol
7DHP	7-Dehydropregnenelone
7TM	7 Transmembrane
AAD	L-Amino acid decarboxylase
AANAT	Arylalkylamine N-acetyltransferase
ACTH	Adrenocorticotrophic hormone
AEA	N-arachidonoyl ethanolamide, anandamide
AFMK	N1-acetyl-N2-formyl-5-methoxykynuramine
AMK	N1-acetyl-5-methoxykynuramine
AVP	Arginine vasopressin
B	Corticosterone
CaR	Calcium receptor
CB1/2	Endocannabinoid receptor 1/2
CGRP	Calcitonin gene-related peptide
cis/trans-UCA	cis/trans-Urocanic acid
CNS	Central nervous system
COMT	Catechol-methyl transferase
COR	Cortisol
CORT	Corticosterone
CRE	cAMP response element
CREB	cAMP response element-binding
CRF/CRH	Corticotropin-releasing factor/hormone
CRF1or 2	Corticotropin-releasing factor receptor 1 or 2
CYP	Cytochrome P

---

DAG	Diacylglycerol
DHEA	Dehydroepiandrosterone
DHT	Dihydrotestosterone
DOC	Deoxycorticosterone
DOR	Delta opioid receptor
DRG	Dorsal root ganglia
DYN A	Dynorphin A
ECD	Extracellular domain
ECS	Endocannabinoids
END	Endorphins
ENK	Enkephalins
ERK	Extracellular signal-related kinase
F	Cortisol
FAAH	Fatty acidamid hydrolase
GABA	Gamma-aminobutyric acid
GI	Gastrointestinal tract
GIRK	G-protein-regulated inwardly rectifying potassium channels
GM-CSF	Granulocyte-macrophage colony-stimulating factor
GPCRs	G-protein-coupled receptors
HaCaT	Human immortalized keratinocytes
HIOMT	Hydroxyindole- <i>O</i> -methyltransferase
HPA axis	Hypothalamic-pituitary-adrenal axis
HPT axis	Hypothalamic-pituitary-thyroid axis
HSD	Hydroxysteroid dehydrogenase
IL	Interleukin
IP3	Inositol trisphosphate
KOR	Kappa-opioid receptor
L-DOPA/DOPA	L-3,4-Dihydroxyphenylalanine
LENK	Leu-Enkephalin
MAO	Monoamine oxidase
MAPK	Mitogen-activated protein kinases
MC2-R	Melanocortin receptor type 2
MEK	Mitogen-activated protein and extracellular signal-regulated kinase
MENK	Met-Enkephalin
MOR	Mu-opioid receptor
MSH	Melanocyte-stimulating hormone
MT1 or 2	Melatonin receptor type 1 or 2
NAS	<i>N</i> -acetylserotonin
NAT	Arylamine <i>N</i> -acetyltransferase
NF- $\kappa$ B	Nuclear factor kappa-light-chain-enhancer of activated B cells
NGF	Nerve growth factor
NO	Nitric oxide

---

NOS	Nitric oxide synthase
NQO1 or 2	Guinone oxidoreductase 1 or 2
OR	Opioid receptor
P450 <sub>scc</sub>	Cytochrome P450 side-chain cleavage enzyme
PDYN	Prodynorphin
PEA	<i>N</i> -palmitoylethanolamide
PENK	Proenkephalin
PH	Phenylalanine hydroxylase
PKA	Protein kinase A
PLC	Phospholipase C
POMC	Proopiomelanocortin
PVN	Paraventricular nucleus
ROS	Radical oxygen species
SP	Substance P
Src	v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog (avian)
SRC	Steroid receptor coactivator
T3	Triiodothyronine
T4	Thyroxine
TGFβ-2	Transforming growth factor beta-2
TH	Tyrosine hydroxylase
THC	Tetrahydrocannabinol
TPH	Tryptophan hydroxylase
TRH	Thyroid-releasing hormone
TRH-R	TRH receptor
TRpOH	5-Hydroxytryptophan
TRPV	Transient receptor potential vanilloid
TRα or β	Thyroid hormone receptor α or β
TSH	Thyroid-stimulating hormone
Tyr	Tyrosinase
URC	Urocortin
Urc 1-3	Urocortin types 1-3
UV	Ultraviolet
UVA	Ultraviolet A radiation
UVB	Ultraviolet B radiation
UVR	Ultraviolet radiation
VDR	Vitamin D3 receptor





# Chapter 1

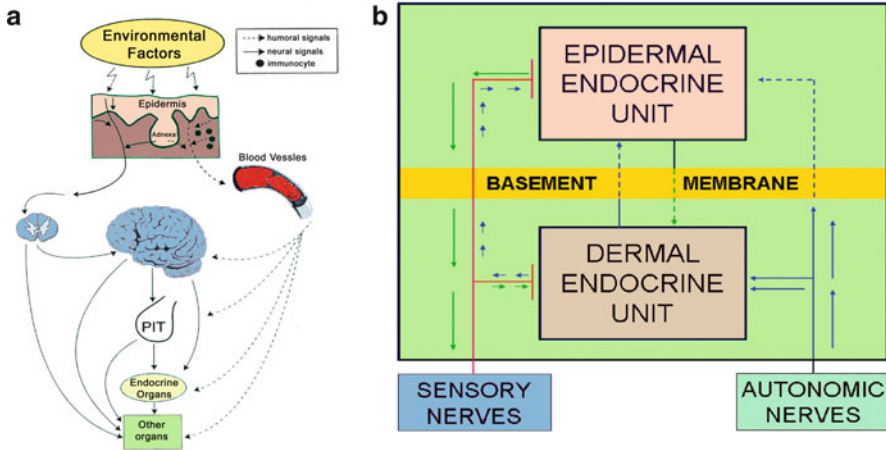
## Introduction

### 1.1 General Overview

The strategic location of the skin as the barrier between the environment and internal milieu determines its critical function in the preservation of body homeostasis, and ultimately organism survival (Slominski 2005; Slominski and Wortsman 2000; Slominski et al. 2000c; Zmijewski and Slominski 2011). It also exposes skin to numerous pathological agents, processes, and events. Thus, the capability to locally recognize, discriminate, and integrate various signals within a highly heterogeneous environment, and to immediately launch appropriate responses, is a vital property of skin (Slominski and Wortsman 2000). These skin functions are integrated into the skin immune, pigmentary, epidermal, and adnexal systems, and are in continuous communication with the systemic immune, neural, and endocrine systems (Arck et al. 2006; Slominski 2009c; Slominski and Wortsman 2000; Slominski et al. 2004c, 2007a; Stenn and Paus 2001).

These fundamental functions result from the location of the skin, which is the largest body organ, at the interphase between external and internal environment, requiring development of efficient sensory and effector capabilities to differentially react to changes in external environment. They are represented by inducible production of biologically active compounds (hormones, neurohormones, and neurotransmitters) that act both locally and at the systemic levels (Fig. 1.1).

The skin being continuously exposed to many external biological or environmental factors (acute transfers of solar, thermal, or chemical energy) had to evolve optimal mechanism(s) to protect, restore, or maintain local and global homeostasis in relation to hostile environment (Slominski et al. 1993b, 2000c; Slominski and Pawelek 1998; Slominski and Wortsman 2000). We have proposed that precise coordination and execution of these responses are mediated by a cutaneous neuroendocrine system, which also is able to reset the body homeostatic adaptation mechanisms (Slominski and Wortsman 2000). Superimposed on this is the impact of psychological stress on skin physiology and pathology,

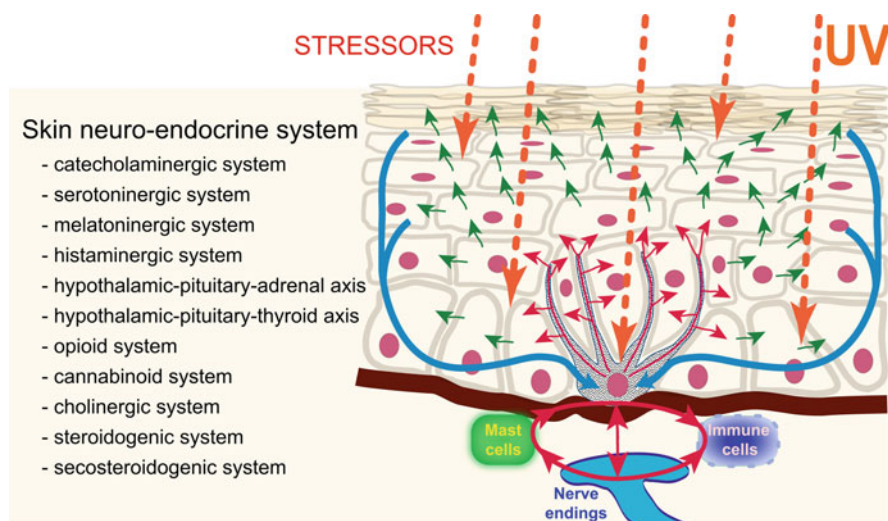


**Fig. 1.1** Skin senses changes in the environment through cutaneous neuroendocrine system, which computes and translates the received information into chemical, physical, and biological messengers that regulate global (A and B) and local (B) homeostasis. These signals travel via through humoral, immune, or neural pathways to reach the central nervous, endocrine, and immune systems as well as other organs. Reproduced with permission from the Endocrine Society (Slominski and Wortsman 2000)

placed in the context of the bidirectional brain–skin communication (Arck et al. 2006; Slominski 2005; Slominski et al. 2008b). To summarize, in reaction to changing external and also internal environment, the skin can generate signals to produce rapid (neural) or slow (humoral or immune) responses at the local and systemic levels (Fig. 1.1).

Coordination between these local and systemic responses is mediated by the skin neuroendocrine system (Slominski and Wortsman 2000), which employs local equivalents of the hypothalamo–pituitary–adrenal axis (HPA) (Slominski et al. 2007a), hypothalamo–pituitary–thyroid (HPT) axis (Pisarchik and Slominski 2002; van Beek et al. 2008), catecholaminergic (Schallreuter et al. 1997), serotonergic, melatoninergic (Slominski et al. 2005c, 2008a), cholinergic (Grando 2006; Grando et al. 2006), steroidogenic (Slominski et al. 2008b), and secosteroidogenic (Bikle 2010; Holick 2003; Slominski et al. 2010) systems (Fig. 1.2). Given their common embryonic origins, it is not surprising that skin shares numerous mediators with the CNS and endocrine system. Recent research has revealed that skin also harbors complex opioidogenic (Grando et al. 1995; Slominski et al. 2011c) and cannabinoidogenic (Biro et al. 2009) systems, the role of which in the maintenance of cutaneous homeostasis is currently being intensively explored.

In this monograph, we will discuss the role of various components of the skin neuroendocrine system in sensing the environment with subsequent regulation of local and global homeostasis with a main focus on the algorithms of classical neuroendocrine axes.



**Fig. 1.2** Skin neuroendocrine system follows the algorithms of classical neuroendocrine or endocrine systems. It also forms a natural platform of signal exchange between internal organs and environment. For this purpose, skin cells not only are subjected to neurohormonal regulation but also do produce neuropeptides, biogenic amines, melatonin, opioids, cannabinoids, acetylcholine, steroids, secosteroids, as well as growth factors and cytokines. Skin neuroendocrine system also entrains immune cells to act as cellular messengers at distal sites

## 1.2 An Overview of Histology and Anatomy

Since histology and anatomy of the skin has been extensively reviewed in three major dermatology and dermatopathology textbooks (Bolognia et al. 2008; Fitzpatrick et al. 1993; Weedon 2010), below we provide only a short overview. The most external layer of the skin, the epidermis, is derived from the ectoderm, and is characterized by a constant renewal. The main constituents of the epidermis, keratinocytes, are either self-replicating in the basal layer (about 50% of basal layer keratinocytes are in this state) or differentiating toward the surface (another 50%). The whole process of differentiation lasts about 31 days. The keratinocytes of succeeding layers (spinous and granular) gradually flatten to form a solid cornified layer that is subsequently shed (this takes another 14 days on average). The intermediate filaments, cytokeratins, are the most important structural elements of the keratinocyte. In the epidermis, cytokeratins 5 and 14 are main cytokeratins in basal keratinocytes and cytokeratins 1 and 10 in differentiating ones. The cornified layer is formed by various cross-linked proteins and lipids. Apart from keratinocytes, there are other cells in the epidermis whose function is more regulatory than structural. Examples are melanocytes, derived from neural crest, which reside in the basal layer. Their density varies in different parts of the body from 1 in 4 to 10 basal keratinocytes. Melanin, protective pigment produced by these cells, is transferred from melanocytes through their processes to approximately neighboring

36 keratinocytes (to form epidermal-melanin unit) by the process of apocoptation. Melanin not only absorbs UV radiation but also serves as a scavenger of reactive oxygen species and miscellaneous chemical compounds. The Langerhans cells are derived from the bone marrow. They reside at different levels of the epidermis and engulf foreign antigens. They transport them to the lymph nodes and present in the context of MHC antigens to T lymphocytes initiating the adaptive immune response.

The dermis is derived from mesoderm. Its bulk is composed of collagen and elastic fibers and glycosaminoglycans. The main collagen of reticular dermis is collagen type I. Collagen type III is present in the adventitial dermis (papillary and peri-appendageal). Elastic fibers are arranged in a parallel manner in the superficial dermis including elaunin fibers (made of microfibrils with elastin core) and perpendicular manner in the papillary dermis (oxytalan fibers made of microfibrils only). Collagen gives skin its strength, elastic fibers its elasticity (ability to retract), and glycosaminoglycans its substance. Various inflammatory cells typically reside in the dermis and increase in numbers when need arises. Dermal vasculature forms superficial and deep dermal plexuses that are connected by straight collaterals. Superficial plexus sends papillary loops toward the surface. Of note, the epidermis does not have its own vasculature and is being nourished through exchange of substances provided by the most superficial parts of papillary capillaries. Glomus bodies (Succquet-Hoyer canals) are important for local thermoregulation.

Skin appendages are of epidermal origin. The hairs cover most of the body. Terminal and vellus hairs differ in their size and function. The hairs undergo cyclic changes of growth (anagen, about 90% of scalp hair, lasts 3–10 years), involution (catagen, 1%, lasts weeks), and rest (telogen, 10%, lasts few months). Of note, different hairs on the body, even directly neighboring, are in different phases of the growth cycle. This is a major difference between humans and animals that shed hair cyclically. The sebaceous glands are usually associated with hair and secrete protective lipid substances by a holocrine mechanism. The coiled eccrine glands are located in the subcutis; their straight ducts transverse the dermis and end in coiled fashion in the acrosyringia of the epidermis. The primary sweat is hyper- or isotonic and becomes hypotonic during passage through the excretory ducts. Sweat production is the most important thermoregulatory mechanism in humans. Apocrine glands are distributed only in some areas of the body (axillae, genital, ear, and eyelid) and have probably only vestigial function in humans.

Last, but not least, the subcutaneous fat tissue is a third important layer of the skin. Fat lobules forming it are separated by fibrous septae transverse rich in vasculature. The adipose tissue is mostly of white type and has important function in isolation, cushion, and energy storage. Often quoted to be body's largest immune/endocrine organ (about 15% of body weight and average surface of about 2 m<sup>2</sup>), skin is a source of multiple mediators and cytokines that act not only locally but also systemically. On the other hand, components of skin respond to internal stimuli and mediators preserving body homeostasis and appropriate functioning.

Skin is studied by a variety of methods. The classic histological slides, prepared from formalin-fixed tissue and stained with hematoxylin and eosin paired with

various special stains and by immunohistochemical methods, are the tools of both dermatopathologist and researcher. Direct immunofluorescence is a complementary method used for both diagnosis and research. Frozen sections are stained here with antibodies against immunoglobulins, complement, and fibrinogen. Different patterns are observed and yield diagnostic information. Popular research tools are the ex vivo skin cell cultures. Both primary (with definite number of cell divisions) and continuous (indefinite number of cell divisions) cell cultures are being used. To better model the conditions present at the skin as tissue, the ex vivo organ cultures are also used. A plethora of cell and molecular biology methods have been applied for studies of both cell and organ cultures. Some popular examples are Western blot, PCR, confocal microscopy, and gene microarrays.

### 1.3 An Overview of Skin Innervation

The skin extensive neural network represented by somatosensory and autonomic nerve fibers has been described in detail in several reviews and books (Bologna et al. 2008; Fitzpatrick et al. 1993; Roosterman et al. 2006; Siemionow et al. 2011; Slominski and Wortsman 2000; Weedon 2010; Yosipovitch 2010). Therefore, below we provide only a short overview.

In the skin, receptors localized on the primary afferent nerve terminals transduce various sensory stimuli, generated upon changes in temperature, pH, and the presence of inflammatory mediators, and convey them to the specific areas of the CNS what results in the perception of pain, itching neuroinflammation, as well as somatic responses of other organs and tissues. The perikarya of cutaneous sensory fibers are localized either in the dorsal root ganglia (DRG) or, those innervating the face and upper neck, in the trigeminal ganglion. Both unmyelinated (C) and myelinated (A) fibers of unipolar sensory cells conduct thresholds at 0.5–2 m/s and 4–70 m/s, respectively. The ortho- and antidromic conduction of afferent nerve fibers results in simultaneous signal transduction and release of neurotransmitters (mainly substance P and CGRP) at the same site. The sensory axons make synapses in dorsal spinal cord neurons depending on somatotopic map of the part of the body surface innervated by the relevant spinal segments. The major ascending routes for sensory cutaneous inputs are via the dorsal column nucleus (DCN) or lateral cervical nucleus (LCN). Both of them transmit to the thalamus, which is a coordinator station for sensory imputes receiving and sending neural signals to somatosensory cortex, midbrain, and hypothalamus—the headquarters of the autonomic nervous system. The connection between thalamus and hypothalamic paraventricular nuclei constitutes important element joining cutaneous stimuli with centers which control body homeostasis and endocrine system, including HPA axis. Also, cutaneous afferent stimuli from face run in the trigeminal root and upon switch in trigeminal nucleus terminate in the thalamus (Siemionow et al. 2011).

The cutaneous innervation has traditionally been considered to consist of a plexus of fibers in the reticular layer of dermis and a more superficial plexus in

the papillary layer, with the majority of sensory endings located in the subpapillary dermis. Recent advances in immunohistochemistry provided an evidence for the existence of intraepidermal nerve fibers (reviewed in Bologna et al. 2008; Legat and Wolf 2009; Roosterman et al. 2006; Slominski and Wortsman 2000; Waller et al. 2011). Intraepidermal nerve terminals associated with Merkel cells, cold receptors, and high-threshold mechanoreceptors have been identified in the basal layer of the epidermis. Thin nerve fibers travel through the dermis, extend into epidermis, and terminate with or without branching in all layers of epidermis including stratum corneum. Waller et al. 2011). The density of epidermal nerve fibers changes during aging and in many pathological conditions like diabetes, psoriasis, or upon ultraviolet radiation. Therefore, quantification of the epidermal nerve fibers' density was proposed to be a valuable prognostic marker for the evaluation of the disease progress (Fromy et al. 2010; Legat and Wolf 2009; Roosterman et al. 2006; Waller et al. 2011).

In the skin, cutaneous nerve fibers have principally sensory character, with an additional component of autonomic nerve fibers distributed exclusively in dermis. Most of them are found in the mid-dermis and the papillary dermis. The autonomic nerves supply arterioles, glomus bodies, hair erector muscles, and apocrine and eccrine glands. A rich network of autonomic and sensory nerve fibers surrounds especially hair follicles, pilosebaceous units, and eccrine and apocrine glands. The sensory and autonomic networks show regional differences according to anatomic location and also have topographical specificity by distributing into well-defined areas called dermatomes. The autonomic nerve fibers in the skin predominantly derive from sympathetic (cholinergic, catecholaminergic, and non-adrenergic/non-cholinergic) and, in the face, rarely parasympathetic (cholinergic) neurons.

In addition to the classic neurotransmitters like acetylcholine, noradrenaline, and serotonin, the postganglionic autonomic nerves in the skin predominantly release also neuropeptides (neuropeptide Y, galanin, vasoactive intestinal peptide, and  $\beta$ -endorphin) and biologically active substances (nitric oxide, ECS) which act as co-transmitters. These compounds modulate the release and activity of the main neurotransmitters and also directly affect targeted cells. Neuropeptides released from cutaneous nerves via a paracrine, juxtacrine, or endocrine manner act on target cells which express specific receptors that are appropriately coupled to an intracellular signal transduction pathway or ion channels, which, when activated, may result in the activation of biological responses such as erythema, edema, hyperthermia, and pruritus.

## Chapter 2

# Biogenic Amines in the Skin

### 2.1 An Overview

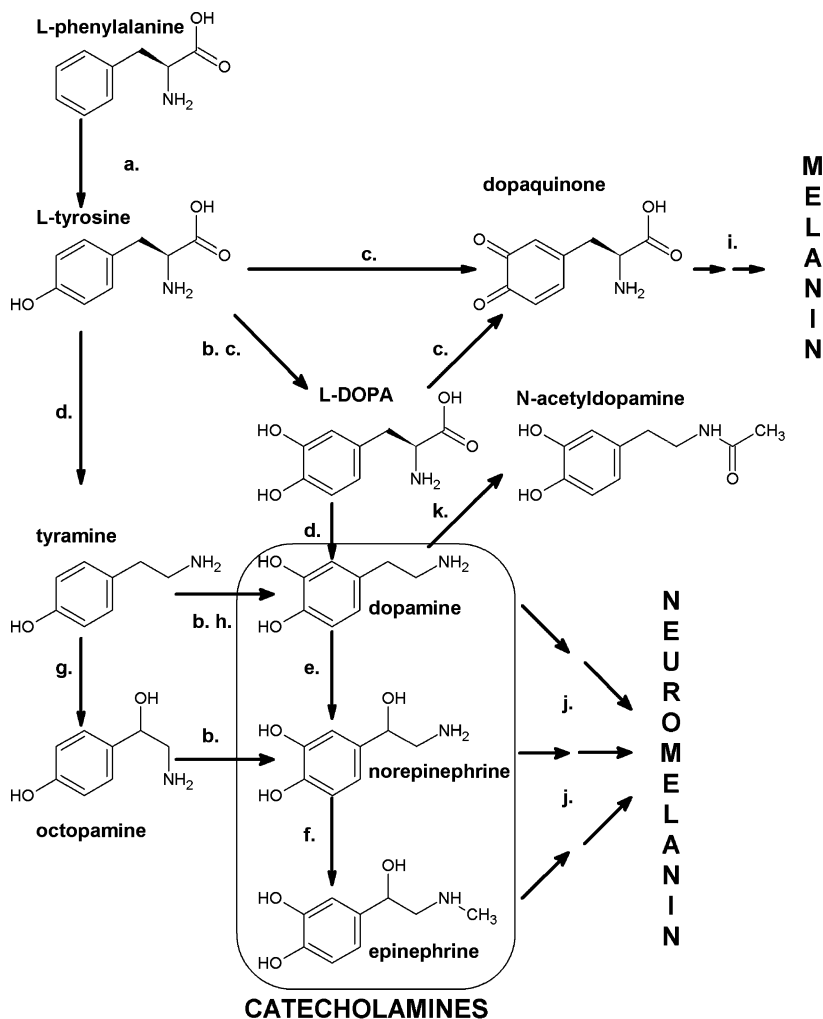
It has been documented that skin resident cells can produce and further metabolize catecholamines, serotonin, and histamine (Fitzpatrick et al. 1993; Gillbro et al. 2004; Schallreuter et al. 1995; Slominski et al. 2005c). These biogenic amines not only regulate phenotype of skin cells cultured in vitro but also can affect skin functions and may have systemic effects (Schallreuter et al. 1997; Slominski and Wortsman 2000; Slominski et al. 2005c). The functional activity of biogenic amines in the skin is mediated through the interactions with specific cell surface receptors (Gillbro et al. 2004; Nordlind et al. 2008; Slominski et al. 2003d); however, non-receptor effects are also considered.

### 2.2 Catecholamines

#### 2.2.1 Production and Metabolism

Nonessential aromatic amino acid L-tyrosine, depending on the cell type and enzymatic context, serves as a direct precursor to catecholamines, tyramine/octopamine (Yen 2001), and melanin pigment (Slominski et al. 2004c). To serve these diverse functions, L-tyrosine is either delivered through the gastrointestinal tract (GI) or produced through phenylalanine hydroxylase (PH)-mediated hydroxylation of L-phenylalanine (Blau et al. 2010; Schallreuter et al. 2008b). L-tyrosine is hydroxylated to L-dihydroxyphenylalanine (L-DOPA) by either tyrosine hydroxylase (TH) or tyrosinase (Tyr), or decarboxylated to tyramine by L-amino acid decarboxylase (AAD) (Fig. 2.1). L-DOPA is further decarboxylated to dopamine by AAD with subsequent hydroxylation and methylation reactions to generate norepinephrine or epinephrine, all of them being oxidated by monoamine oxidase (MAO) or methylated by catechol-methyl transferase (COMT) (Fig. 2.1). L-DOPA





- a. L-Phenylalanine hydroxylase
- b. Tyrosine hydroxylase
- c. Tyrosinase
- d. L-amino acid decarboxylase
- e. Dopamine  $\beta$ -monooxygenase

- f. Phenylethanolamine N-methyltransferase
- g. Dopamine  $\beta$ -hydroxylase
- h. Cytochrome P450 (Cyp2D)
- i. Melanogenesis
- j. Neuromelanogenesis
- k. Acetylation

**Fig. 2.1** Catecholamine synthesis in the skin. The common pathway in the skin requires its consecutive hydroxylations of L-phenylalanine [mediated by phenylalanine hydroxylase (PH)] to L-tyrosine with following hydroxylation by tyrosine hydroxylase (TH) or tyrosinase to produce L-dihydroxyphenylalanine (L-DOPA). L-DOPA is either oxidized to DOPA quinone with following multistep transformation to melanin or serves as a substrate for synthesis of catecholamines. The skin expresses complete enzymatic machinery required for dopamine synthesis (L-amino acid

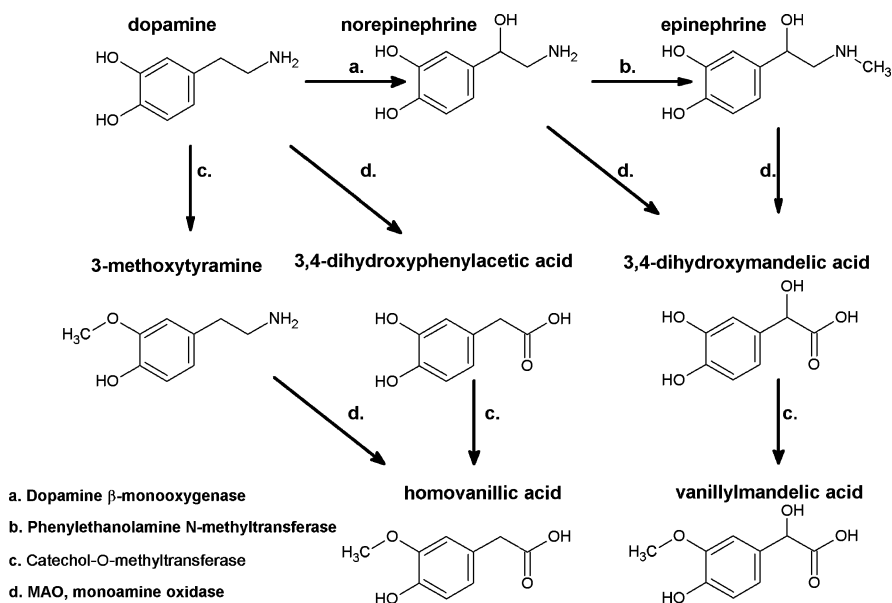
and catecholamines can also be oxidized by either tyrosinase or metal cations to form melanin and neuromelanin pigments (Fitzsimons et al. 2002; Lassalle et al. 2003; Park et al. 2009; Slominski et al. 2004c) (Fig. 2.1).

Human epidermal keratinocytes and melanocytes have the capability to synthesize the catecholamines from L-tyrosine with sequential production of L-DOPA, dopamine, norepinephrine, and epinephrine through the action of classical enzymes listed above with the subsequent inactivation of catecholamines by MAO or COMT (Fig. 2.2) (Fujiwara et al. 2005; Gillbro et al. 2004; Schallreuter et al. 1992, 1995). Interestingly, acetylation of dopamine to *N*-acetylDOPA has also been described in the hamster skin (Gaudet et al. 1993). Activity of TH and PH depends on local availability of their essential cofactor/electron donor, i.e., 6R-L-erythro-5, 6, 7, 8-tetrahydrobiopterin (6BH4) as demonstrated for the first time by Schallreuter's group (Schallreuter et al. 1994, 1997). Importantly, Schallreuter and coworkers demonstrated de novo synthesis/recycling/regulation of 6BH4 in both human epidermal keratinocytes and melanocytes as well as in hair follicles (Schallreuter et al. 1997, 1998). Furthermore, AAD activity requires pyridoxal phosphate (PP) as the cofactor, the cutaneous availability of which is regulated locally (Coburn et al. 2003). Lymphocytes and other immune cells can also represent an additional source of catecholamines: L-DOPA production with its further transformation to epinephrine and norepinephrine has been shown in human lymphocytes (Musso et al. 1997) as well as in Langerhans cells (Falck et al. 2004). An additional cutaneous source of catecholamines is their dermal release from adrenergic nerve fibers (Fitzpatrick et al. 1993). A challenging task in current skin biology is to determine which skin cells and adnexal structures have similar capability of de novo synthesis of catecholamines and what is the final product in different compartments.

An important alternative source of L-DOPA for cutaneous catecholamines is its production via the tyrosine hydroxylase activity of tyrosinase that, depending on the intracellular environment including acidic pH, may not undergo oxidation but will diffuse or be transported to other cells or systemic circulation (Slominski et al. 2004c, 2011a). In fact, diffuse "melanocytic organ" can provide DOPA or its adducts to systemic circulation to serve either as a precursor for further modifications or as a bioregulator (Slominski et al. 1993a, 2011a; Zmijewski and Slominski 2009a). A role for tyrosinase-derived L-DOPA is supported by findings that retinal network adaptation to bright light requires tyrosinase-dependent production of DOPA (Page-McCaw et al. 2004). This phenomenon represents the TH-independent pathway of peripheral dopamine synthesis (Eisenhofer et al. 2003) and

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**Fig. 2.1** (continued) decarboxylase, AAD) and its subsequent conversion into norepinephrine (dopamine  $\beta$ -hydroxylase) and methylation (phenylethanolamine *N*-methyltransferase) to form epinephrine. An alternative pathway of catecholamine synthesis involves decarboxylation of L-tyrosine to tyramine, which in turn is hydroxylated by TH (and Cyp2D) or dopamine  $\beta$ -hydroxylase to octopamine or dopamine, respectively. Octopamine could be metabolized to norepinephrine by TH. This alternative pathway which is present in invertebrates remains to be tested in the skin. Catecholamines also undergo oxidation to corresponding quinoinones with further multistep transformation to neuromelanin, a process similar to melanogenesis starting from L-DOPA



**Fig. 2.2** Catecholamine catabolism. Catecholamines are deactivated by L-monoamine oxidase (MAO) and Catechol-*O*-methyltransferase (COMT) leading to the synthesis of homovanillic acid (from dopamine) or vanillylmandelic acid from norepinephrine or epinephrine. Alternatively, as shown for dopamine metabolism the order of reaction may be changes with COMT acting first and then followed by MAO

it can regulate activities of melanocytes and immune cells (Slominski and Paus 1990; Slominski et al. 1998c). These findings are in agreement with our hypothesis that L-tyrosine and L-DOPA can have hormone- and neurotransmitter-like roles (Slominski and Paus 1990, 1994; Slominski et al. 2011a), with melanocytes acting as important regulators of catecholamines' availability in the skin (Slominski et al. 1993a).

## 2.2.2 Bioregulatory Role of Catecholamines in the Skin

### 2.2.2.1 Dopamine Receptors

There are five subtypes of dopamine receptors, and they have been categorized into two families, i.e., D1-like receptors (D1 and D5) and D2-like receptors (D2, D3, and D4) (Watson 1994). The D1-like receptor agonists stimulate G<sub>s</sub>-dependent intracellular production of cAMP (Missale et al. 1998). The D2-like receptor agonists activate G<sub>i</sub> proteins and inhibit intracellular cAMP signaling pathway (Missale et al. 1998; Watson 1994). In addition, via Gβγ subunits, D2-like receptors are capable of inhibiting N- and L-type calcium channels which results in the

activation of G-protein-regulated inwardly rectifying potassium channels (GIRKs) (Beaulieu and Gainetdinov 2011). After D2-like receptors were identified in the keratinocytes (Fuziwara et al. 2005) they were found to play a significant role in the maintenance of epidermal barrier homeostasis. Application of D2-like receptor agonists accelerated barrier recovery, whereas D2-like receptor antagonists delayed it. Actual receptor subtypes localize to different parts of the epidermis: D4 is localized in the uppermost part of the epidermis and D2 is localized in the basal layer of the epidermis where it plays a role in the regulation of cell proliferation (Fuziwara et al. 2005). It remains to be tested whether dopamine is also regulating epidermal and follicular pigmentary systems as well as adnexal functions including hair follicle.

Dopamine receptors on lymphocytes exert differential effects. Dopaminergic signaling through D2-like receptors of T lymphocytes showed an immunostimulatory effect (Besser et al. 2005), whereas signaling through D1-like receptors had immunoinhibitory effect (Saha et al. 2001). Dopamine also inhibits proliferation of human lymphocytes and causes apoptosis of peripheral blood mononuclear cells (Bergquist et al. 1997). IL-6 (and other cytokines) stimulates a development of a subtype of T lymphocytes capable of producing IL-17 (and other cytokines), i.e., Th17 lymphocytes. Th17 lymphocytes constitute relatively recently described branch of immune responses (Harrington et al. 2006). Dopamine released by dendritic cells induces IL-6–Th17 axis and upregulates synovial inflammation (Nakano et al. 2011). The IL-6–Th17 axis plays a role in the pathogenesis of inflammatory diseases including rheumatoid arthritis. It can therefore be deduced that dopamine may also have various differential modulatory roles in the skin immune system.

### 2.2.2.2 Adrenergic Receptors

The adrenergic receptors belong to the classic seven-transmembrane G-protein-coupled receptor (GPCR) family. These receptors respond to catecholamines and can be subdivided into subtypes of  $\alpha$  and  $\beta$  families, based on their differential pharmacological responses and protein sequences (Lands et al. 1967). More specifically, these receptors are defined, in part, by their endogenous ligand affinity to  $\beta$  receptors having a higher affinity to epinephrine when compared to norepinephrine, and to  $\alpha$  receptors having a higher affinity for norepinephrine. Alpha adrenergic receptors can be further subdivided into  $\alpha_1$  and  $\alpha_2$ , and  $\beta$  receptors can be further subdivided into  $\beta_1$ ,  $\beta_2$ , and  $\beta_3$  subtypes. The  $\alpha_1$  ( $\alpha_{1a}$ ,  $\alpha_{1b}$ , and  $\alpha_{1d}$ ) receptors couple to phospholipase C via  $G_{q\alpha}$  and stimulate the formation of diacylglycerol and inositol trisphosphate (Cotecchia 2010). The  $\alpha_2$  ( $\alpha_{2a}$ ,  $\alpha_{2b}$ , and  $\alpha_{2c}$ ) receptors couple to  $G_{i\alpha}$  and inhibit the formation of cAMP, whereas  $\beta$  receptors are positively coupled to the formation of cAMP via  $G_{s\alpha}$  (Hein 2006).

Various receptors of both  $\alpha$  and  $\beta$  subfamilies of adrenergic receptors are present on epidermal and dermal cells (Grando et al. 2006; Schallreuter et al. 1995). As expected,  $\alpha$  and  $\beta$  receptors are also expressed in dermal blood vessels. Their

activation by catecholamines causes vasoconstriction and decreases vascular permeability (Ding et al. 1995; Harada et al. 1996).

Keratinocytes express mainly  $\beta_2$  receptors and also  $\alpha_1$  receptors (Steinkraus et al. 1992; Drummond et al. 1996; Sivamani et al. 2007). Stimulation of  $\beta$ -adrenergic receptors in epidermal keratinocytes results in increased cAMP production, activation of protein kinase C, and formation of inositol-1,4,5-trisphosphate, calcium influx, and extracellular signal-related kinase (ERK) dephosphorylation through the action of serine/threonine phosphatase PP2A (Chen et al. 2002; Pullar et al. 2001; Schallreuter et al. 1995). Catecholamines stimulate keratinocyte differentiation with increased expression of keratins 1, 10, involucrin, and transglutaminase (Mammone et al. 1998; Schallreuter et al. 1995). Moreover, there is a local gradient of receptor expression with the highest level in basal keratinocytes and decreasing level toward the surface of the epidermis (Schallreuter et al. 1995). This indicates a potential stimulatory functional role of catecholamines in the process of keratinocytes' differentiation. Catecholamine- $\beta_2$  adrenergic system has been implicated in the pathogenesis of atopic dermatitis, psoriasis, and vitiligo (Sivamani et al. 2007). Expression of  $\beta_2$  receptors is increased in vitiligo and decreased in psoriasis (Schallreuter et al. 1993; Takahashi et al. 1996). In vitiligo, there is an overproduction of 6-BH4 leading to a dysregulation of catecholamine biosynthesis with increased plasma and epidermal norepinephrine levels. This is associated with high numbers of  $\beta_2$  adrenoceptors in differentiating keratinocytes and with a defective calcium uptake in both keratinocytes and melanocytes (Schallreuter et al. 2008a). In atopic eczema, a point mutation in the beta 2-adrenoceptor gene could alter the structure and function of the receptor, thereby leading to a low density of receptors on both keratinocytes and peripheral blood lymphocytes (Schallreuter et al. 1997). It is also known that catecholamines and  $\beta$  receptors have a role in wound healing although their exact role is far from being clarified (Ghoghawala et al. 2008; Pullar et al. 2008) (see also discussion of fibroblast below). The adrenergic beta-receptors not only affect keratinocytes' proliferation and differentiation but also their immune activities. Activation of  $\beta$  receptors on keratinocytes affects expression of  $\beta$ -defensin 3 (Martin-Ezquerria et al. 2011).

Studies on cultured melanoma cell lines have shown that catecholamines can be an additional factor affecting melanogenesis (Howe et al. 1991). Their role in the function of the pigmentary system has been well described in nonhuman systems (reviewed by Slominski et al. 2004c). Human melanocytes express  $\alpha_1$  and  $\beta_2$  receptors (Gillbro et al. 2004; Hu 2000; Hu et al. 2000; Scarparo et al. 2000; Schallreuter et al. 1996). Activation of  $\alpha_1$  receptors leads to the IP3-DAG signaling (Schallreuter et al. 1996) and  $\beta_2$  receptor activation leads to cAMP signaling (Gillbro et al. 2004).  $\beta_2$  but not  $\alpha_1$  receptor activation induces pigmentation (Gillbro et al. 2004; Schallreuter et al. 1996). The expression of  $\beta_2$  receptors on human melanocytes increases in response to UV irradiation (Yang et al. 2006). UVB irradiation increases epinephrine release by cultured keratinocytes that in turn increases pigmentation in co-cultured melanocytes, which is an example of the interactions between these two cell types (Sivamani et al. 2009).

Adrenergic receptors are expressed also on immune cells of the dermis (Steinkraus et al. 1996). Binding of adrenergic agonists to their receptors on lymphocytes has immunostimulatory effect and affects their homing. On the contrary, stimulation of  $\beta$  receptor usually has immunosuppressive effects, but in other model systems can also cause immunostimulation, i.e., increase the number of lymphocytes (Bergmann and Sautner 2002).

Mouse Langerhans cells express  $\alpha 1$ ,  $\beta 1$ , and  $\beta 2$  adrenergic receptors (Seiffert et al. 2002), and it was shown that epinephrine and norepinephrine inhibit the ability of Langerhans cells to present antigens (Seiffert et al. 2002).

Agonists of  $\beta 2$  receptors on mast cells inhibit the release of preformed mediators such as histamine, and also newly synthesized mediators such as prostaglandin D2 from mast cells (Okayama and Church 1992). They also inhibit release of inflammatory cytokines from mast cells (Bissonnette and Befus 1997).  $\beta$  receptors are expressed on dermal fibroblasts (Pullar and Isseroff 2006; Pullar et al. 2008). Ligation of  $\beta 2$  receptors activates epidermal growth factor (EGF) receptor and extracellular signal-regulated kinase (ERK) signaling that in turn stimulates fibroblast migration. Binding of agonists to the  $\beta 2$  receptors can also activate protein A kinase (PKA) which can stimulate cell proliferation (Pullar and Isseroff 2006), attenuate collagen gel contraction, and alter actin cytoskeleton and focal adhesion distribution via PKA-dependent mechanisms (Pullar and Isseroff 2006). A link between body stress response system that results in the release of epinephrine and activation of intracellular signaling that leads to DNA damage has been shown recently (Hara et al. 2011). Specifically, in mouse and human fibroblasts binding of agonists to the  $\beta 2$  receptors led to Gs-protein-dependent activation of protein kinase A, followed by the recruitment of beta-arrestins. Then,  $\beta$ -arrestin 1 facilitated AKT-mediated activation of MDM2 and also promoted MDM2 protein binding to and degradation of p53 protein by acting as a molecular scaffold. The degradation of p53 resulted in the lack of protection and DNA damage (Hara et al. 2011).

### 2.2.2.3 Non-receptor-Mediated Effects of Catecholamines

In the skin there are several potential non-receptor-mediated effects, which are based on autooxidation of catecholamines in alkaline environment with a velocity increased by metal cations (Lassalle et al. 2003; Slominski et al. 2004c). The potential phenotypic implications are predominantly based on the well-documented activity of L-DOPA which through its oxidation products and active melanogenesis can affect functions of immune cells (Slominski and Goodman-Snitkoff 1992; Slominski et al. 2009b). The possible mechanisms of action were discussed previously (Slominski et al. 1998c, 2004c) and, therefore, have been shortly summarized below. L-DOPA dramatically inhibits an in vitro phosphorylation of glycoproteins dependent on the presence of Mn ions indicating action of quinones generated through oxidation of DOPA (Slominski and Friedrich 1992). It can also affect cellular metabolism in melanotic cells (Scislawski et al. 1984, 1985). Also, diffusible products of DOPA oxidation are potent inhibitors of lipid peroxidation

(Memoli et al. 1997), and 5-S-cysteinyl-dopa inhibits hydroxylation/oxidation reactions induced by the Fenton reaction (Napolitano et al. 1996). The potential cycling from indole to quinone forms of L-DOPA and its derivatives may affect levels of reactive oxygen/nitrogen species or oxidation of intracellular proteins and lipids (Tsang and Chung 2009). Finally, both free and protein-bound L-DOPA can trigger expression of several antioxidant enzymes including superoxide dismutase or NAD(P)H:Quinone oxidoreductase (NQO1) (Nelson et al. 2007). Thus, taking into consideration similar chemical properties of DOPA and catecholamines (products of DOPA enzymatic metabolism), and that their oxidation leads to the production of neuromelanin, one can safely conclude that non-receptor-mediated effects and mechanisms will be similar to that described for DOPA (Slominski et al. 2011a). Taking into consideration the above chemical properties of dopamine, epinephrine, or norepinephrine, one can expect that at micromolar or higher concentrations the predominant effects will be non-receptor-mediated mainly through their oxidation products and neuromelanin polymers generated during this process. It is also possible that some of the phenotypic effects at lower concentrations could also be influenced by oxidative effects.

#### **2.2.2.4 Conclusions**

Dopamine, epinephrine, and norepinephrine are produced in the skin resident and nonresident cells. Their phenotypic effects are mediated through activation of dopaminergic and adrenergic receptors, the expression of which is cell-type and cell-context dependent. Their roles in epidermal, dermal, and adnexal as well as skin immune functions remain to be further investigated. There are also non-receptor-mediated mechanisms shared by their precursor, L-DOPA. It is likely that cutaneous catecholaminergic system will communicate with brain by activating sensory nerves, or, with other tissues, via entry into systemic circulation and by affecting immune cells circulating from the skin to other organs (Fig. 1.1).

### **2.3 Histamine**

#### ***2.3.1 Production and Metabolism of Histamine***

Histamine is derived from the decarboxylation of histidine by the L-histidine decarboxylase. After release, histamine is degraded by histamine-N-methyltransferase (in brain and at periphery) or diamine oxidase (in the periphery) (Fitzpatrick et al. 1993; Zhang et al. 2007). Histamine is produced mainly by mast cells and basophils. Cross-linking of IgE antibodies attached to the cell membrane represents a main mechanism for histamine release. Histamine binds to four different types of seven-transmembrane receptors that signal through G-proteins.

The H<sub>1</sub> receptor is found on smooth muscle and endothelial cells and is responsible for smooth muscle contraction and decreased adhesion of endothelial cells. H<sub>2</sub> receptor is located on vascular smooth muscles and parietal cells in the stomach and is responsible for vasodilatation and gastric acid secretion. H<sub>3</sub> receptor is found in the central and peripheral nervous systems and is responsible for decreased secretion of several neurotransmitters including histamine, acetylcholine, serotonin, and norepinephrine. H<sub>4</sub> receptor is found primarily on basophils and has a role in chemotaxis (Fitzpatrick et al. 1993; Zhang et al. 2007).

### ***2.3.2 Bioregulatory Role of Histamine in the Skin***

In the epidermis, H<sub>1</sub> and H<sub>2</sub> receptors are expressed on keratinocytes (Albanesi et al. 1998; Koizumi and Ohkawara 1999; Koizumi et al. 1998; Shinoda et al. 1998) and H<sub>2</sub> receptors on epidermal melanocytes (Yoshida et al. 2000). Mediators released from mast cells inhibit keratinocyte growth in culture (Huttunen et al. 2001). Activation of keratinocyte H<sub>2</sub> receptors affects proliferation and differentiation via activation of the cyclic AMP pathway and also phospholipase C pathway with associated increase in intracellular calcium levels (Koizumi and Ohkawara 1999). In mouse keratinocytes, H<sub>2</sub> receptor signaling through the PLC second messenger system is inhibited during calcium-induced keratinocyte differentiation by an autocrine loop which involves downregulation of H<sub>2</sub> receptor expression and inhibition of histamine metabolism (Fitzsimons et al. 2002). In keratinocytes, activation of the H<sub>1</sub> receptor enhances UVB-induced IL-6 production (Koizumi and Ohkawara 1999; Koizumi et al. 1998), whereas H<sub>1</sub> receptor antagonists inhibit ICAM-1 expression (Ling et al. 2004). Histamine upregulates keratinocyte MMP-9 production via the H<sub>1</sub> receptor (Gschwandtner et al. 2008). H<sub>2</sub>, however, not H<sub>1</sub>, agonists stimulate intracellular calcium signaling in keratinocytes (Koizumi and Ohkawara 1999). In these cells, histamine acting on H<sub>1</sub> receptors increases the expression of IFN- $\gamma$ -induced intercellular adhesion molecule 1 (ICAM-1) and MHC class I molecules. It also augments IFN- $\gamma$ -induced release of chemokines such as CXCL10, as well as the release of GM-CSF via protein kinase C $\alpha$  and extracellular signal-regulated (ERK) kinase (Giustizieri et al. 2004; Kanda and Watanabe 2004). In cultured keratinocytes, histamine through the activation of H<sub>1</sub> receptor inhibits CCL17 production by suppressing p38 MAP kinase and NF- $\kappa$ B activities. Histamine acts as a negative feedback signal for existing Th2-dominant inflammation by suppressing CCL17 and enhancing CXCL10 production (Fujimoto et al. 2011). The effect of histamine acting through H<sub>2</sub> receptor appears to be the opposite. Histamine, via H<sub>2</sub> receptor, increases survival of keratinocytes acting by NF- $\kappa$ B activation (Kim and Lee 2010). IL-17, produced by Th17 cells infiltrating into the dermis (a cytokine involved in various inflammatory skin diseases including psoriasis), stimulates keratinocytes to produce inflammatory mediators such as IL-36, TNF- $\alpha$ , IL-6, and IL-8 (Carrier et al. 2011). Histamine markedly augments the production of IL-8 and GM-CSF in the presence of IL-17 and TNF- $\alpha$  in



keratinocytes (Moniaga et al. 2011). Moreover, histamine induces human  $\beta$ -defensin 2 and 3 production in keratinocytes acting via  $H_1$  receptors by activating NF- $\kappa$ B, AP-1 pathway, or STAT1, STAT3, and AP-1 as well as JAK2 and MEK/ERK signaling pathways (Ishikawa et al. 2009; Kanda and Watanabe 2007). Histamine promotes cutaneous antimicrobial defenses and wound repair by stimulating secretion of defensins (Ishikawa et al. 2009; Kanda and Watanabe 2007). Histamine also enhances nerve growth factor production by inducing c-Fos expression in keratinocytes (Kanda and Watanabe 2003).

The activation of the  $H_2$  receptors on melanocytes stimulates melanogenesis (Yoshida et al. 2000). Histamine, similarly to  $\alpha$ -MSH, contributes to hyperpigmentation by enhancing eumelanin/pheomelanin ratio (Lassalle et al. 2003). Acting at the  $H_2$  receptor histamine stimulates melanocyte migration in culture via signaling through ERK, CREB, and Akt (Kim and Lee 2010). Histaminergic system is upregulated in the B16F10 melanoma cells when compared to noncancerous melanocytes, which indicates that it might have a role in tumorigenesis (Davis et al. 2011). Both Western blot and immunohistochemical studies showed much stronger histidine decarboxylase expression in melanoma cells as compared to normal melanocytes (Haak-Frendscho et al. 2000). Moreover,  $H_1$  histamine receptor antagonists were shown to induce genotoxic and caspase-2-dependent apoptosis in human melanoma cells, but not normal melanocytes (Jangi et al. 2006).

In the dermis, histamine receptors are expressed on fibroblasts, immunocytes, endothelial cells, blood vessels, smooth muscle, and nerve endings (Fitzpatrick et al. 1993). In Th2 lymphocytes stimulation of  $H_4$  receptor led to the activation of transcription factor AP-1 followed by the release of IL-31, which is involved in the development of pruritus (Gutzmer et al. 2009). On the other hand, activation of  $H_4$  histamine receptors expressed on monocytes activated intracellular calcium mobilization and inhibited the CCL2 chemokine production which reduced recruitment of monocytes (Dijkstra et al. 2007). Histamine acts on  $H_4$  receptors of eosinophils and mediates their chemotaxis, induces cell shape change, and upregulates adhesion molecules CD11b/CD18 (Mac-1) and CD54 (ICAM-1). This effect, while observed in cultured eosinophils, may be of paramount importance in the skin (Ling et al. 2004).

Histamine also acts on  $H_2$  and  $H_4$  receptors of plasmacytoid dendritic cells and downregulates production of TNF- $\alpha$ , IFN- $\alpha$ , and CXCL8 (Mazzoni et al. 2003). Plasmacytoid dendritic cells migrate in response to  $H_4$  receptor agonist stimulation. Of note,  $H_4$  receptor is present in high levels on plasmacytoid dendritic cells in the lesional psoriatic skin (Gschwandtner et al. 2011).

### 2.3.3 Conclusions

Histamine is produced not only by mast cells but also by other cells of epidermis and dermis and acts locally in the epidermis and dermis by binding to  $H_1$ - $H_4$  receptors. Histamine targets not only endothelium and smooth muscles of blood

vessels but also modulates function of keratinocytes, melanocytes, and cells of skin immune system. It affects intracellular signaling cascades, cell proliferation, and melanogenesis. Histamine is upregulated in melanoma cells. It signals mainly via  $H_4$  receptor on the cells of the immune system and affects their migration and cytokine secretion patterns. Moreover, it modulates Th2-type immune responses and antimicrobial peptide expression. Thus, histamine is an important part of the neuro-immuno-endocrine system of the skin (Slominski and Wortsman 2000) with local and systemic effects (Figs. 1.1 and 1.2).

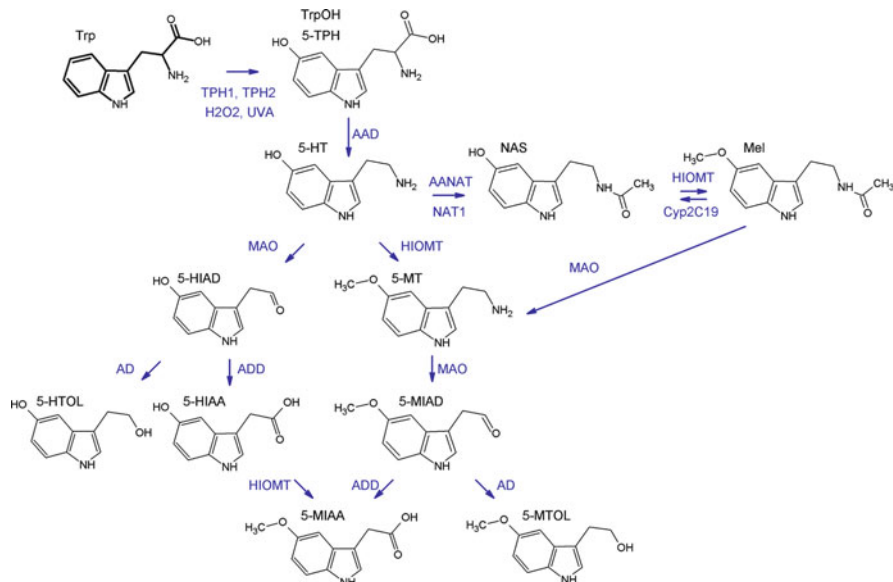
## 2.4 Serotonergic System

### 2.4.1 *Production and Metabolism of Serotonin*

#### 2.4.1.1 An Overview

Serotonin (5-hydroxytryptamine, 5-HT) is widely synthesized throughout the animal kingdom, plants, and unicellular organisms (Azmitia 2001, 2007). In plants, serotonin serves as a trophic factor and an antioxidant which is similar to the animal kingdom (Azmitia 2001). In humans, serotonin was shown to be synthesized predominantly by intestinal enterochromaffin cells with other sites of production represented by the central nervous system, pineal gland, retina, ovaries, placenta, thymus, pancreas, skin, breast, gestational tissues, blood vessels, rectal epithelium, bronchial epithelial cells, thyroid parafollicular cells, mast cells, and lymphocytes (Nordlind et al. 2008).

The first obligatory step in the synthesis of serotonin is the hydroxylation of L-tryptophan to produce 5-hydroxytryptophan (TrpOH) in a reaction catalyzed by tryptophan hydroxylase (TPH) (Mockus and Vrana 1998), a protein encoded by either TPH1 gene expressed ubiquitously (Mockus and Vrana 1998) or TPH2 gene expressed predominantly in the brain (Zhang et al. 2004). This reaction requires oxygen and cofactor 6BH4. TrpOH is further decarboxylated by AAD to produce 5-HT. In humans, L-tryptophan is present in blood plasma at steady-state level both in the free form (approximately  $1.2 \times 10^{-5}$  M) and bound to serum albumins (ca.  $6 \times 10^{-5}$  M), with TPH having a  $K_m$  for tryptophan of approximately  $10^{-8}$  M. Thus, fluctuations in free pool of tryptophan directly and immediately alter the level of serotonin synthesis (Nordlind et al. 2008). Catabolism of serotonin is initiated by MAO with the production of 5-hydroxyindoleacetaldehyde, oxidized further by aldehyde dehydrogenase (E.C. 1.2.1.3) to 5-hydroxyindole-3-acetic acid (5-HIAA), which is the main product of metabolism, or reduced to 5-hydroxytryptophol (HTOL) by alcohol dehydrogenase (E.C. 1.1.1.1) (Fig. 2.3). 5-HT can also be methylated to 5-methoxytryptamine (5MTT) and catabolized as shown in Fig. 2.3. Additional pathway involves serotonin acetylation by arylalkylamine *N*-acetyltransferase (AANAT) or arylamine *N*-acetyltransferase isoenzyme showing substrate specificity toward both arylamines and arylalkylamines to produce *N*-acetylserotonin (NAS)



**Fig. 2.3** Biochemical pathway of serotonin synthesis and metabolism in the skin. The pathway starts with hydroxylation of tryptophan by tryptophan hydroxylase type 1 or 2 (TPH1 or TPH2) to form 5-hydroxytryptophan (5-TPH; TrpOH). TrpOH can also be produced by nonenzymatic action of UVA and H<sub>2</sub>O<sub>2</sub>. Serotonin (5-hydroxytryptamine, 5-HT) derives from 5-TPH by action of L-amino acid decarboxylase, AAD. Serotonin can be acetylated by aralkylamine *N*-acetyltransferase (AANAT) or *N*-acetyltransferase (NAT) to produce *N*-acetylserotonin (NAS) with further methylation by hydroxy-indole-*O*-methyl transferase (HIOMT) to melatonin. Deactivation of serotonin is catalyzed mainly by MAO with the formation of 5-hydroxyindoleacetaldehyde (5-HIAD) which is followed by the action of alcohol (AD) or aldehyde dehydrogenase (ADD) with the formation of 5-hydroxytryptophanol (5-HTOL) or 5-hydroxyindole-3-acetic acid (5-HIAA), respectively. Alternatively, HIOMT activity may also lead to the production of methylated derivatives of serotonin. The first step catalyzed by HIOMT leads to the formation of 5-methoxytryptamine 5-MT. The subsequent action of MAO results in 5-methoxyindoleacetaldehyde (5-MIAD) formation. Finally, AD or ADD facilitates the synthesis of 5-methoxytryptophol (5-MTOL) or 5-methoxyindole-3-acetic acid (5-MIAA), respectively. HIOMT was found also to catalyze the conversion of 5-HIAA to 5-MIAA. By the action of MAO melatonin can be metabolized to 5-methoxytryptamine (5-MT), thus entering the pathway leading to 5-MTOL or 5-MIAA formation

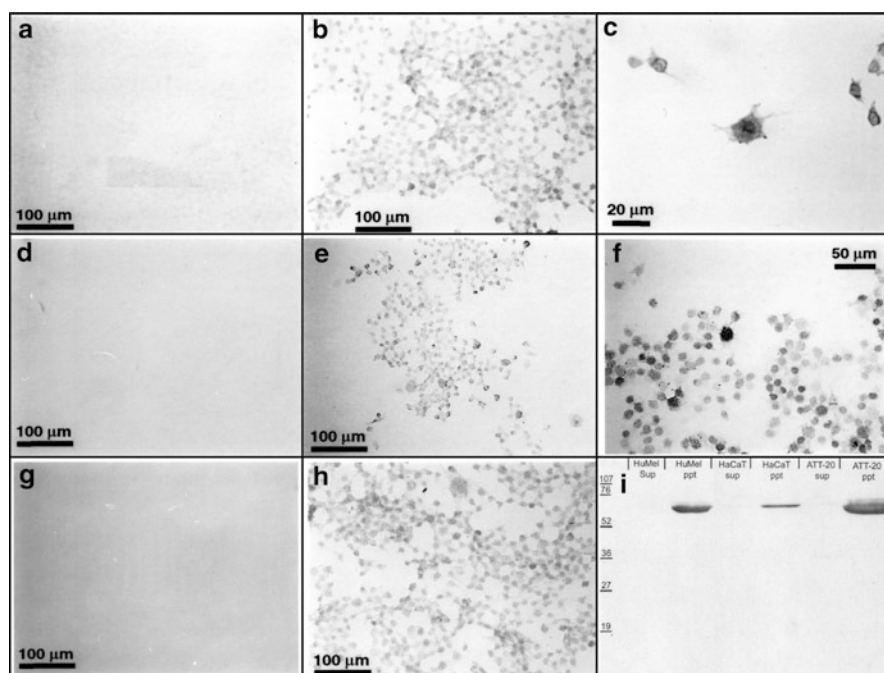
(Fitzsimons et al. 2002; Klein 2004). NAS can also be further metabolized to melatonin (Reiter 1991). In the skin, a number of NAS metabolites unrelated to melatonin were found, the nature and mechanism of generation of which remain to be defined (Slominski et al. 2003b, c). After release into blood, serotonin is actively taken up into platelets and stored in solid granules with the help of a serotonin transporter (5-HTT), a member of the Na<sup>+</sup>/Cl<sup>-</sup>-dependent transporter superfamily, which actively regulates serotonin transport. Serotonin can be transported through the plasma membrane in either direction; however, under most conditions, its reuptake is favored (Nordlind et al. 2008). Plasma serotonin is also cleared by the liver and lung endothelial cells and further catabolized to 5-HIAA.

### 2.4.1.2 Production and Metabolism of Serotonin in the Skin

Mammalian skin cells can produce serotonin via the sequential transformation of L-tryptophan by TPH and AAD (Slominski et al. 2005c) (Fig. 2.3). Thus, the *TPH1* gene is expressed in human skin under normal and pathological conditions as well as in a wide array of normal and transformed human epidermal, dermal, and adnexal skin cells with some cells expressing the aberrant *TPH1* transcript (Slominski et al. 2002c, 2003b, c). As to the *TPH2* gene, it is expressed in the retinal pigment epithelium (Zmijewski et al. 2009b) and normal and malignant melanocytes (Zmijewski and Slominski, unpublished). Although the *TPH* gene is expressed almost in all types of human skin cells, the highest expression was found in normal and malignant melanocytes that also accumulated significant amounts of serotonin (Figs. 2.4 and 2.5) (Slominski et al. 2003a, 2005c). Interestingly, the enzymatic conversion of tryptophan to TrpOH in melanoma cells occurs at high levels, comparable to those in the brain (Slominski et al. 2002a, c). TPH and *TPH1* were also detected in the mouse and hamster skin, and in cultured mouse follicular melanocytes and melanoma cells (Slominski et al. 2002a, 2003b, c). Interestingly, the *TPH1* gene expression changes during murine hair cycle (Slominski et al. 2003b, c). In addition, TPH and serotonin are strongly expressed in rodent mast cells. It is also important to notice that the skin has a capability for de novo synthesis/recycling of the 6BH4 (Schallreuter et al. 1997, 1998, 2008a) and of pyridoxal 5'-phosphate (PLP) (Coburn et al. 2003) both serving as important cofactors necessary for the production of TrpOH and serotonin. Interestingly, nonenzymatic production of TrpOH through H<sub>2</sub>O<sub>2</sub> and UVA radiation indicates that a free-radical-mediated oxidation of L-tryptophan is also possible in the skin (Schallreuter et al. 2008a).

In human skin biopsies immunoreactivity of TPH and serotonin was found in normal epidermal melanocytes and malignant melanomas (Figs. 2.4 and 2.5) (Slominski et al. 2003a) with additional detection by immunofluorescence techniques in epidermal keratinocytes, hair follicles, eccrine glands, blood vessels, and skin mast cells (Slominski et al. 2005c). These findings are consistent with the immunodetection of serotonin in perivascular human mast cells of adrenal cortex (Lefebvre et al. 2001) and breast epithelial cells (Matsuda et al. 2004). Serotonin was also detected by immunocytochemistry in dermal Merkel cells in rat and pig skin at the epidermal rete ridges and upper hair follicles adjacent to nerve terminals (Nordlind et al. 2008). Cutaneous serotonin content can be affected by inflammatory processes (Lonne-Rahm et al. 2008; Nordlind et al. 2008; Rasul et al. 2011; Thorslund et al. 2009). For example, human skin affected by psoriasis or chronic eczema showed elevated expression of serotonin in the epidermal and adnexal structures (Nordlind et al. 2008).

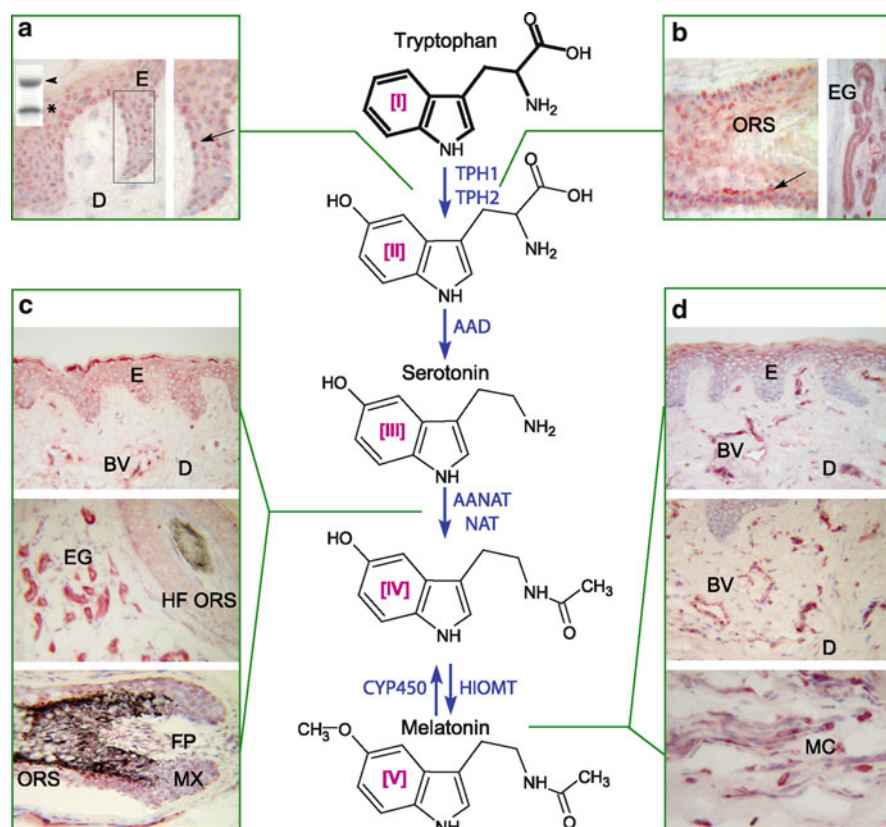
The catabolism of serotonin in mouse skin is initiated by its deamination by MAO, followed by the oxidation or reduction of the resultant 5-hydroxyindole acetaldehyde to 5-HIAA and/or 5-HTPOL (Slominski et al. 2003b, c). Similar metabolism was uncovered in rat skin, although in this species 5-HIAA was the



**Fig. 2.4** Expression of TPH, serotonin (5-HT), and serotonin transporter (5-HTT) in skin cells. Panels **a–h** show immunocytochemical detection of 5-HT (**b, c**), 5-HTT (**e, f**), and TPH (**h**) in fixed cells using corresponding antibody at the dilution of 1:5,000 (antibody against 5-HT, Diasporin Corp., Stillwater, MN) or 1:1,000 (antibodies against TPH and 5-HTT, Chemicon, Temecula, CA). (**a, d, g**) Negative controls incubated with secondary antibody only. (**i**) Western blot showing detection of 5-HTT in membranous (ppt) but not cytosolic (sup) fractions from human melanoma (HuMel), HaCaT keratinocytes (HaCaT), and ATT-20 pituitary cells. For technical details of the assays, see Slominski et al. (2005d)

main degradation product and 5-HTPOL remained below the limit of detectability (Semak et al. 2004). MAO metabolism of serotonin was also detected in guinea pig skin (Tachibana et al. 1990) and production of 5-HIAA was documented in human epidermal keratinocytes and melanoma cells (Slominski et al. 2002c).

The alternative serotonin metabolism pathway in the skin is represented by its acetylation to *N*-acetylserotonin, which in human and rodent skin and cultured skin cells is mediated via the action of either AANAT or NAT with mixed arylamine/arylalkylamine substrate specificity (Slominski et al. 2005c). In hamster skin, we characterized two *N*-acetyltransferase activities including NAT-1 with substrate specificity toward arylamines, and NAT-2 showing substrate specificity toward both arylamines and arylalkylamines such as serotonin, tryptamine, and methoxytryptamine (Gaudet et al. 1993; Slominski et al. 2002a). Furthermore, we demonstrated that at least part of this activity in hamster, rat, and human skin represented native AANAT (Slominski et al. 2002a). In accordance, serotonin *N*-acetyltransferase activity was significantly inhibited by low concentrations of



**Fig. 2.5** Melanonegic system in the skin. TPH1 Western blot insert in the Panel **a** is of approximately 50 kDa (arrowhead) that is processed and/or degraded to lower molecular weight species (asterisk). It was immunolocalized in the epidermis (ES), hair follicle (ORS), eccrine glands (EG), showing the highest expression in melanocytes (arrows) (Panels **a** and **b**). 5-hydroxytryptophan is further decarboxylated by aromatic amino acid decarboxylase (AAD). AANAT (enzyme acetylating serotonin) is expressed in cells of epidermal, dermal, and adnexal compartments (E, BV, EG, and hair follicle structures in Panel **c** on the left). Immunocytochemical localization of melatonin-like immunoreactivity is shown in Panel **d** on the right (upper E, BV, and MC). Immunocytochemistry was performed on human skin biopsies: *E* epidermis, *D* dermis, *BV* blood vessel, *EG* eccrine gland, *HF ORS* hair follicle outer root sheath, *FP* hair follicle papilla; *MX* hair follicle matrix, *MC* mast cells. For technical details, see Slominski et al. (2005d). Reproduced with permission from the publisher (Slominski et al. 2008a)

coenzyme A-S-N-acetyltryptamine [Cole bisubstrate; BSI, see (Hickman et al. 1999; Khalil et al. 1998)], indicating true AANAT activity. However, significant enzymatic activity generating NAS was resistant to BSI suppression, showing that in rodents arylamine activity (NAT-2) resistant to BSI can also participate in the acetylation of serotonin (Semak et al. 2004; Slominski et al. 2002a). Rodent NAT-2 is a homologue of human NAT-1; thus, it is likely that NAT-1 may contribute to NAS production also in the human skin. Interestingly, in the C57BL/6 mouse

producing inactive AANAT (Roseboom et al. 1998), we detected cutaneous transformation of serotonin to NAS and, to a lesser extent, acetylation of tryptamine (Slominski et al. 2003b, c). Most interestingly, acetylation of serotonin, but not of tryptamine, was dependent on the phase of hair cycle, skin anatomic location, and the presence of pathology (melanoma). NAS was further metabolized to several products (the chemical nature of which remains to be defined) in a hair cycle-dependent fashion (Slominski et al. 2003b, c). In humans, both skin racial pigmentation and cutaneous pathology determine the reaction rate and specificity of serotonin acetylation (Slominski et al. 2002c).

### **2.4.2 Bioregulatory Role of Serotonin in the Skin**

Serotonin regulates a wide range of physiological processes at the central and peripheral levels acting as a neurotransmitter, hormone, cytokine, biological modifier, growth factor, morphogen, and antioxidant or pro-oxidant (Azmitia 2007, 2010). The above functions are mediated through receptor-dependent and receptor-independent mechanisms (Hoyer et al. 2002).

Serotonin acts via multiple receptor subtypes labeled as 5-HT1 through 5-HT7 (Hoyer et al. 2002). Most of these receptors are metabotropic, with the exception of 5-HT3, which is ionotropic and primarily gates sodium and potassium ions. 5-HT1 receptors (1A, 1B, 1D, 1E, and 1F) couple via  $G_{i\alpha}$  to inhibit cAMP formation while 5-HT4, 5-HT6, and 5-HT7 all couple via  $G_{s\alpha}$  to stimulate cAMP production (Hoyer et al. 2002). In addition, 5-HT1A receptors produce membrane hyperpolarization by coupling to  $K^+$ -channels. 5-HT2 (2A, 2B, and 2C) receptors couple via  $G_{q\alpha}$  to phosphatidylinositol hydrolysis and the formation of inositol trisphosphate and diacylglycerol (Hoyer et al. 2002). The 5-HT5 receptor (5A and 5B) is considered to be an orphan receptor. Serotonin receptor function can be modulated by RNA editing, endogenous lipids that act as allosteric modulators, and serotonin moduline (tetrapeptide, 5-HT-moduline) that is produced by proteolytic modification of chromogranin. 5-HT moduline is an allosteric modulator which regulates 5-HT5 receptor dimerization and formation of either homodimers or heterodimers. The receptors' heterogeneity and functional diversity are also amplified by the process of alternative splicing and differential subunit incorporation into the receptor complex. The regulation of 5-HT receptor activity is also affected by serotonin transporters, which remove serotonin from the extracellular environment or, under certain conditions, pump it out of the cell.

In human skin and skin cells, we identified expression of genes coding 5-HT receptors, including *HTR1A*, *1B*, *2A*, *2B*, *2C*, and *7* genes, and it was shown that the pattern of expression was cell type specific and modified by skin pathology (Slominski et al. 2003d). Interestingly, alternatively spliced form of *HTR2C* with a deletion of exon 2, a fragment of exon 3, and an insertion of cryptic exon containing termination codon was found in human melanoma, while the *HTR2B* isoform with a deletion of exon 2, but with a preserved reading frame coding for a receptor protein without transmembrane domains 3 and 4 was found in normal

human skin and skin affected by basal cell carcinoma (Slominski et al. 2003a). We also found RNA editing (A to G substitution) in human *HTR7* gene (Slominski et al. 2003a), which may be connected to the local expression of adenosine deaminases. In mouse and hamster skin, expression of the *HTR2B* and *HTR7* genes was demonstrated, which was dependent on the phase of hair cycle (mouse) and type of tissue or cells (Slominski et al. 2004b).

We should also mention that Kaneko et al. have failed to detect 5-HT<sub>2A</sub> gene in epidermal keratinocytes (Kaneko et al. 2009). However, these findings have to be considered with caution, since other researchers demonstrated that 5-HT<sub>2A</sub> antagonists inhibited UVR-induced skin carcinogenesis (Sreevidya et al. 2008, 2010) and that sunlight-induced immunosuppression could be mediated via the activation of 5-HT<sub>2A</sub> by cis-urocanic acid (Walterscheid et al. 2006). Furthermore, 5-HT<sub>2A</sub> protein was detected by immunocytochemistry in dermal lymphocytes, fibrocytes, vasculature, and sensory nerve endings, abating the epidermis (Nordlind et al. 2008), while 5-HT<sub>1A</sub> receptor was localized to keratinocytes of the upper epidermis, epidermal melanocytes, mast cells, and dermal vasculature (Nordlind et al. 2008). Furthermore, 5-HT<sub>1A</sub> and 5-HT<sub>2A</sub> were detected in the majority of benign tumors such as compound nevi, dysplastic nevi, and also in malignant melanomas (Nordlind et al. 2008). By the use of immunocytochemistry, 5-HT<sub>2C</sub> was detected in epidermal Langerhans cells and melanocytes, 5-HT<sub>3</sub> in the basal epidermal keratinocytes, and 5-HT<sub>7</sub> in dermal vasculature (Nordlind et al. 2008). 5-HT<sub>1A</sub>, 2A, and 2C were also detected in rodent skin dermal and epidermal immune cells (Nordlind et al. 2008). Diverse expression of 5-HT receptors was also found in immune cells that were dependent on cell type and their level of activation.

Also Merkel, Langerhans and mast cells, lymphocytes and macrophages (Nordlind et al. 2008), and immortalized human epidermal keratinocytes and melanoma cells express 5-HTT (Fig. 2.4). Their role is substantiated by observations which showed that serotonin uptake inhibitors could induce spontaneous bruising, pruritus, urticaria, angioedema, erythema multiforme, the Steven–Johnson syndrome, toxic epidermal necrolysis, erythema nodosum, alopecia, hypertrichosis, leukocytoclastic vasculitis, and acneiform eruption (reviewed by Nordlind et al. 2008; Slominski et al. 2005c). This can also be associated with flares of psoriasis vulgaris and development of delayed hypersensitivity.

Under in vitro conditions, serotonin exerted variable effects on skin cells depending on the context (Nordlind et al. 2008; Salim and Ali 2011; Slominski et al. 2005c). It stimulated proliferation of dermal fibroblasts (Slominski et al. 2005c), similarly to non-skin fibroblasts (Seuwen and Pouyssegur 1990). Serotonin also stimulated growth of epidermal melanocytes in the absence of growth factors, while inhibiting their proliferation in media supplemented with serum (Slominski et al. 2003a). The former effect could be linked with the stimulation of intracellular cAMP accumulation, while the latter could represent serotonin antagonism with serum growth factors (Slominski et al. 2005c). NAS, the product of serotonin metabolism, showed no effect on the proliferation of fibroblasts and melanocytes (Slominski et al. 2003a) and serotonin or inhibitors of its uptake inhibited melanogenesis (reviewed by Slominski et al. 2004b; Slominski et al. 2005c). In addition,



serotonin modulated proliferation of cultured murine keratinocytes (Maurer et al. 1997). Interestingly, serotonin content within mast cell granules steadily decreased throughout anagen and increased during catagen and telogen phases of hair cycle (Hasse et al. 2007).

Serotonin shows vasoactive and immunomodulatory effects. For example, it plays a role in the Arthus reaction (Tachibana et al. 1990; Yuasa et al. 2001), induces sustained vascular permeability (Fujii et al. 1994), and also modulates the inflammatory response to substance P (SP) via capsaicin-sensitive sensory fibers (Khalil and Helme 1990). Serotonin participates in the activation of T cells and natural killer cells by macrophages, initiation of delayed-type hypersensitivity responses, production of chemotactic factors, and the modification of innate immune responses (Benton et al. 2010; Betten et al. 2001; Cloez-Tayarani and Changeux 2007; Hsueh et al. 2002; Mossner and Lesch 1998). In allergic contact dermatitis and psoriasis, the number of cells expressing both 5-HT1A and tryptase diminishes, whereas the number of dermal cells expressing 5-HT2A and CD3 increases, including atopic dermatitis (Lonne-Rahm et al. 2008; Nordlind et al. 2008; Rasul et al. 2011; Thorslund et al. 2009). Similar pattern is found in the murine epidermis affected by contact eczema. Furthermore, both eczematous and psoriatic human skin shows increased number of mononuclear cells expressing 5-HTT (reviewed by Nordlind et al. 2008). In addition, serotonin can act as a chemoattractant for eosinophils, probably by binding to 5-HT2A receptors. It is involved in the mast cell recruitment to the site of tissue injury through the activation of 5-HT1A, however, without inducing their degranulation (Nordlind et al. 2008). Regulatory function of 5-HT1A in inflammatory responses is emphasized by the suppression of the severity of contact allergy in rats, after topical or oral administration of its agonist, buspirone (Nordlind et al. 2008). Another 5-HT1A agonist, tandospirone, attenuates itching in patients with atopic dermatitis (Nordlind et al. 2008). On the other hand, treatment with 5-HT2A antagonists reduced the severity of contact allergic reactions in mice and one of them, spiperone, was effective when applied either systemically or topically. Furthermore, 5-HT2 receptor antagonist, ketanserin, inhibited the established but not challenge-induced phases of allergic contact dermatitis (Nordlind et al. 2008). Serotonin is also involved in the pathogenesis of cholestatic and uremic pruritus, urticaria, and itch reaction (reviewed by Slominski et al. 2005c).

### ***2.4.3 Serotonin Receptors on Sensory Nerves***

5-HT receptors were widely detected on cutaneous sensory nerve endings (reviewed by Nordlind et al. 2008; Slominski et al. 2005c). Intradermal injection of serotonin into rat elicited enhanced c-fos-like immunoreactivity in superficial lamina at the lateral aspect of the dorsal horn, in a manner similar to the immunoreactivity evoked by capsaicin. The 5-HT receptor was detected in unmyelinated sensory axons at the dermal-epidermal junction and the nerve endings of Pacinian

corpuscles of rat glabrous skin (Carlton and Coggeshall 1997) and rat sinus hair follicle (Tachibana et al. 2005). 5-HT<sub>1</sub> receptors are present in the dermis of rabbits on afferent nerve fibers around hair follicles and sebaceous glands (Branchek et al. 1988). 5-HT<sub>2A</sub> receptors are partially responsible for mediating scratching in mice (Tachibana et al. 1990). Although neither 5-HT<sub>2</sub> nor 5-HT<sub>3</sub> appears to be involved in itch responses caused by chronic allergic skin dermatitis in rats, acute scratching is mediated by skin 5-HT<sub>2</sub> receptors, and intradermal injection of serotonin induced itching in normal, but not inflamed skin (reviewed by Nordlind et al. 2008; Slominski et al. 2005c). In human skin, 5-HT<sub>2A</sub> and 5-HT<sub>3</sub> are localized on sensory nerve ending in the dermis or located close to or entering the epidermis, and their activation may explain pruritic responses to intradermally injected serotonin (Nordlind et al. 2008; Slominski et al. 2005c). Specifically, an antagonist of 5-HT<sub>3</sub>, ondansetron, can reduce the severity of pruritus, while paroxetine is used in the treatment of pruritus and its antipruritic action is connected with downregulation of 5-HT<sub>3</sub> expression (Nordlind et al. 2008; Slominski et al. 2005c).

#### ***2.4.4 Reception of Ultraviolet Light***

The cutaneous serotonergic system may play a role in body reception of and reaction to light (Slominski et al. 2005c). For example, it has been reported that UVA-induced well-being can be linked to increased serum serotonin and decreased melatonin levels after a single radiation exposure (Gambichler et al. 2002). It has also been proposed that 5-HT<sub>2A</sub> plays a role in the transduction of UVR energy into biological responses by serving as the receptor for cis-urocanic acid (cis-UCA), generated through photoisomerization of the trans-UCA in the stratum corneum after absorption of UVR (Walterscheid et al. 2006). Cis-UCA acts as a powerful local and systemic immunosuppressor (Garssen et al. 2001), and it was proposed that 5-HT<sub>2A</sub> mediates immunosuppressive effects of UVR after binding of cis-UCA (Walterscheid et al. 2006). A role for 5-HT<sub>2A</sub> in UVB-induced skin photocarcinogenesis was also suggested (Sreevidya et al. 2008, 2010). Other authors proposed that cis-UCA and serotonin mediate UVB-induced immunomodulation, however, via independent pathways in which cis-UCA does not act through 5-HT<sub>2A</sub> (Kaneko et al. 2009). Thus, there is sufficient information to support involvement of the local serotonergic system in cutaneous responses to the UV light; however, the mechanism may be more complex than originally anticipated. It may include activation of 5-HT receptor signaling on either nerve ending or skin cells secondary to UVR-induced local production of serotonin or alternative ligands for HT receptors with a consequent regulation of local homeostasis and immune system. Such signals will be projected to the brain via the ascending nerve routes. Furthermore, release of serotonin into circulation may generate endocrine effects.

### **2.4.5 Conclusions**

The mammalian skin cells have the capability to produce and metabolize serotonin. The cutaneous phenotypic effects are mediated by its interactions with 5-HT receptors including 5-HT<sub>1A</sub>, 1B, 2A, 2B, 2C, 3 and 7, and 5-HTT receptors, which are expressed in a cell type-dependent manner. The serotonin receptors are also expressed on sensory nerve endings, which transmit to the brain information on changes in skin homeostasis induced by either intrinsic or environmental factors (Slominski 2005; Slominski and Wortsman 2000). The topical application of specific receptors agonists or antagonists, serotonin uptake inhibitors or modulation of local serotonin production/degradation may represent future novel therapies of skin diseases including neurodermatoses and itching disorders. Finally, the cutaneous serotonergic system may be involved in the transformation of light energy of solar radiation into local and systemic biological responses, with the latter mediated via transmission to brain, endocrine effects, or regulation of systemic responses as shown on Figs. 1.1 and 1.2.

## Chapter 3

# Melatonineric System in the Skin

### 3.1 Melatonin Production

Melatonin production is highly conserved in nature through different species including bacteria, unicellular eukaryotes, algae, plants invertebrates, and vertebrates (Hardeland et al. 2011; Reiter 1991; Slominski et al. 2008a; Tan et al. 2002; Yu and Reiter 1993). In mammals, melatonin is produced in the pineal gland (Reiter 1991) as well as in brain, retina, Harderian gland, ciliary body, lens, thymus, airway epithelium, bone marrow, immune cells, gonads, placenta, gastrointestinal tract, and skin (Bubenik 2002; Carrillo-Vico et al. 2004; Hardeland et al. 2011; Kanda and Watanabe 2007; Pandi-Perumal et al. 2006; Slominski et al. 2005a, 2008a; Watson 1994; Zmijewski et al. 2009b), and perhaps other organs. Circulating melatonin predominantly derives from the pineal gland by diffusion into the circulation, although entry from other extra-pineal sites of production is also possible.

Melatonin is a product of a two-step transformation of serotonin which involves acetylation catalyzed by AANAT to NAS (a rate-limiting step) followed by methylation by hydroxyindole-*O*-methyltransferase (HIOMT, EC 2.1.1.4) to produce melatonin (*N*-acetyl-5-methoxytryptamine) (Reiter 1991; Yu and Reiter 1993). In the pineal gland melatonin production is controlled by the suprachiasmatic nucleus through nocturnal sympathetic release of norepinephrine that acting via adrenergic receptors activates cAMP-dependent signal transduction cascades leading to the stimulation of AANAT and ultimate production of melatonin (Klein 2007; Reiter 1991; Yu and Reiter 1993). Melatonin synthesis is also potentiated by vasoactive intestinal peptide (VIP), pituitary adenylate cyclase-activating peptide (PACAP), and neuropeptide Y (Klein 2007; Reiter 1991; Yu and Reiter 1993). NAS can also be produced by the action of arylamine *N*-acetyltransferases (Fitzsimons et al. 2002) as it was shown in human (Slominski et al. 2002c), rat (Semak et al. 2004), hamster (Gaudet et al. 1993; Slominski et al. 2002a), and murine skin (Slominski et al. 2003b, c). NAS can be further methylated to melatonin, depending on the anatomic location and activity of HIOMT. This is best illustrated in the C57BL/6

mice, defined by some authors as a natural melatonin “knockdown” (Kobayashi et al. 2005; Roseboom et al. 1998; Slominski et al. 2003b, c). Specifically, in the C57BL/6 mouse serotonin can be acetylated to NAS in a reaction mediated by an enzyme different from conventional AANAT providing an important mechanistic explanation for the significant production of melatonin in the peripheral organs of this species, which express HIOMT (Ma et al. 2008; Scarparo et al. 2000; Slominski et al. 2003b, c). In addition, the existence of low flux rate alternative pathways has been proposed that involves *O*-methylation of serotonin with subsequent *N*-acetylation, or *O*-methylation of tryptophan followed by consecutive decarboxylation and *N*-acetylation.

Transcripts of *AANAT* and of *HIOMT* genes were detected in normal and pathological skin biopsies, and in most skin cells cultured in vitro including normal keratinocytes (neonatal and adult, epidermal and follicular), immortalized HaCaT keratinocytes, fibroblasts (dermal and hair follicle papilla), normal melanocytes, several melanoma cell lines, and squamous cell carcinoma cells (Slominski et al. 2002b). Interestingly, novel isoforms of *AANAT* and *HIOMT* were detected in normal and pathological skin (invaded by basal cell carcinoma cells) and in neonatal keratinocytes (Slominski et al. 2002b). Gene expression in epidermal and dermal skin cells was followed by the synthesis of the AANAT and HIOMT enzymes with the detection of corresponding enzymatic activities (Slominski et al. 2002b).

The acetylation of serotonin was also dependent on local cellular environment. Thus, when AANAT activity was calculated for two substrates, tryptamine and serotonin, the activity ratios were close to 1 for all melanoma lines and for keratinocytes. On the other hand, these ratios ranged from 2.5 to 6 for whole skin from three white subjects and zero in melanocytes and in whole skin of a black subject whose AANAT activity toward tryptamine was below detectability level. These findings suggest a role for both skin racial pigmentation and type of cutaneous pathology (such as melanoma) in this regulation (Slominski et al. 2002b). Both of them may be important determinants of reaction rate and specificity of serotonin acetylation. Using immunocytochemistry AANAT antigen was detected in suprabasal differentiating keratinocytes in human scalp epidermis. However, melanocytes also exhibited immunoreactivity for this enzyme (Fig. 2.5). High expression of the antigen was also seen in the outer peripheral epithelial layers of the anagen hair follicles (Fig. 2.5) and the basal cells of the sebaceous and eccrine glands. The expression was further found in sensory nerve endings abutting the epidermal layers (Slominski et al. 2005c). Melatonin-like immunoreactivity in human skin was detected on differentiating keratinocytes in spinous and granular layers of the epidermis (Fig. 2.5). The antigen was not expressed in keratinocytes of basal and suprabasal layers of the epidermis, while being found in singly scattered melanocytes. Melatonin immunoreactivity was also detected throughout the hair follicle epithelium, in blood vessels, and cutaneous mast cells (Slominski et al. 2005c). These findings were further confirmed by the detection of NAS and melatonin using tandem liquid chromatography/mass spectrometry (LC/MS) in epidermal cells (Slominski et al. 2002a, c) and hair follicles (Kobayashi et al.

2005). These findings showed that human skin, in addition to the pineal gland and retina, possesses the intrinsic capability to synthesize melatonin (Abe et al. 1999; Carrillo-Vico et al. 2004; Finocchiaro et al. 1991; Itoh et al. 1999; Scarparo et al. 2000). Importantly, this cutaneous melatonergic pathway operates in a compartment-specific manner since it is localized mainly to the epidermal, adnexal, and dermal cell populations (Fig. 2.5) (Slominski et al. 2008a).

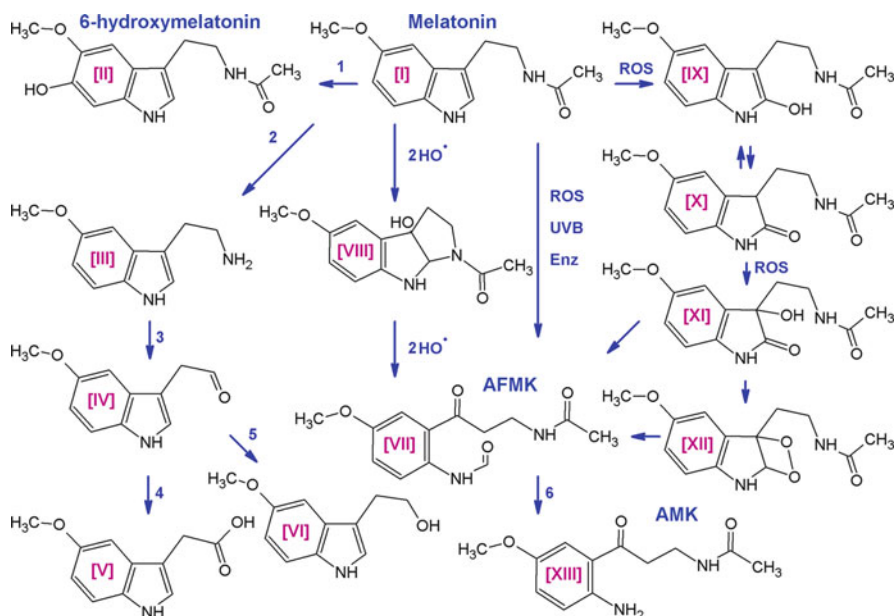
Similar capability to produce melatonin was demonstrated in rodent skin (Slominski et al. 2005c). For example, in hamster skin fragments maintained *ex vivo* serotonin was transformed into melatonin with NAS as the intermediate product (Slominski et al. 1996b, 2002a). This transformation was time- and dose-dependent, and was stimulated by forskolin—indicating involvement of cAMP signal in this process (Slominski et al. 1996b). These findings have been confirmed in follow-up studies (Slominski et al. 2005c). Specifically, biochemical assays in mouse, rat, and hamster skin clearly demonstrated that skin of all of these species can transform serotonin to NAS, the obligatory precursor for melatonin (Semak et al. 2004; Slominski et al. 2002a, 2003b, c). Additionally, murine skin in organ culture and mouse vibrissae hair follicles can produce melatonin and its synthesis was enhanced by the addition of norepinephrine (Kobayashi et al. 2005). Interestingly, detailed analysis with bisubstrate Cole inhibitor in combination with molecular analyses showed that in rodent skin NAS production was initiated by both AANAT and NAT (Semak et al. 2004; Slominski et al. 2002a), while in C57BL/6 mouse NAS appeared to be only produced by NAT (Slominski et al. 2003b, c). This latter finding provides mechanistic explanation for melatonin production in C57BL/6 mice at selected extracranial sites, which would require HIOMT expression since NAS produced via AANAT-independent pathways could serve as substrate for HIOMT-mediated transformation into melatonin (Ma et al. 2008; Scarparo et al. 2000; Slominski et al. 2003b, c). Our enzymatic studies excluded corporal skin of the C57BL/6 mouse *in vivo* as a site of melatonin production, although we detected low levels of HIOMT activity in mouse ear (Slominski et al. 2003b, c).

## 3.2 Melatonin Degradation

Melatonin can be degraded via indolic and kynuric pathways. The first one involves 6-hydroxylation by CYP1A1, CYP1A2, or CYP1B1 to 6-hydroxymelatonin (predominantly in the liver), which after sulfatation or glucuronidation is excreted in urine (Kopin et al. 1961; Ma et al. 2005, 2008). In the liver, the intrinsic clearance for melatonin hydroxylation by high- and low-affinity components indicated that both mitochondrial and microsomal cytochrome P450s metabolize melatonin principally by 6-hydroxylation, with *O*-demethylation representing minor metabolism (Ma et al. 2005). In addition, melatonin deacetylase produces 5-methoxytryptamine that is oxidized by monoamine oxidase to form 5-methoxyindoleacetaldehyde, which is converted to 5-methoxyindole acetic acid by aldehyde dehydrogenase or to 5-methoxytryptophol by alcohol dehydrogenase (Cahill and Besharse 1989; Grace

et al. 1991). In the kynuric pathway, melatonin can be converted either enzymatically or non-enzymatically to N1-acetyl-N2-formyl-5-methoxykynuramine (AFMK), in a process that encompasses generation of 3-hydroxymelatonin, 2-hydroxymelatonin, melatonin 2-indolinone, 3-hydroxymelatonin, 2-indolinone, and melatonin dioxetane as intermediate products (Hardeland et al. 2009; Hirata et al. 1974; Reiter et al. 2007). AFMK synthesis involves enzymes or pseudoenzymes such as cytochrome c, horseradish peroxidase, indoleamine dioxygenase, myeloperoxidase, oxoferryl hemoglobin, or hemin as well as nonenzymatic pathway that may be activated in the presence of reactive oxygen species (ROS) or UVB (Fischer et al. 2006a; Hardeland et al. 2009; Kanda and Watanabe 2007; Seever and Hardeland 2008; Semak et al. 2005, 2008). In addition, catalase, arylamine formamidase, hemoperoxidase, and ROS can stimulate the conversion of AFMK to AMK (Hardeland et al. 2009; Kanda and Watanabe 2007; Reiter et al. 2007). Melatonin can also be demethylated to NAS by CYP2C19 or CYP1A2 (Semak et al. 2008). However, according to some authors, AFMK and AMK pathways of melatonin metabolism are insignificant at the systemic level in mouse (Ma et al. 2008).

Melatonin metabolites 5-methoxytryptamine (5-MTT) and 5-methoxytryptophol (5-MTOL) have been detected in cultured mammalian skin fragments and melanoma cells (Slominski et al. 1996b, 2002b, c), indicating similarity in the degradative pathways of melatonin metabolism in frog skin and retina (Cahill and Besharse 1989; Grace et al. 1991), including the activity of monoamine oxidase (MAO) in mammalian skin (Semak et al. 2004; Slominski 2005). It was shown that cutaneous degradation of melatonin may also include pathways known to be operative in the liver and kidney (Grace et al. 1991; Kanda and Watanabe 2007; Pandi-Perumal et al. 2006) with 6-hydroxymelatonin production as an intermediate (Fischer et al. 2006a). This shows that indolic degradative pathway is operating in the skin (Fischer et al. 2006a; Slominski et al. 1996b). However, experiments with cultured human immortalized keratinocytes have shown that melatonin is mostly metabolized to 2-hydroxymelatonin and AFMK by the kynuric pathway or through direct non-enzymatic action of UVB (Fischer et al. 2006a). Interestingly, UVB also induces AFMK utilization by keratinocytes, suggesting the involvement of arylamine formamidase in the further metabolism of AFMK to AMK (Fischer et al. 2006a). Based on the above, together with the known mechanism for melatonin degradation or transformation in peripheral organs, we proposed that in the skin melatonin can be metabolized via alternative pathways including nonenzymatic reactions (Fig. 3.1). These would exhibit species-, site-, tissue-, and cell compartment- as well as cell type-dependent differences, subjected to further modulation by environmental factors including UVR (Fischer et al. 2008b; Slominski et al. 2008a). Pathways' activities and the nature of the final product would be linked to the spatial distribution of melatonin in the skin, to the specific cell type and subcellular compartments, where the biological activity of melatonin would either be attenuated due to its degradation or be amplified by the generation of even more potent metabolites, such as AFMK or AMK (Fischer et al. 2008b; Slominski et al. 2008a).

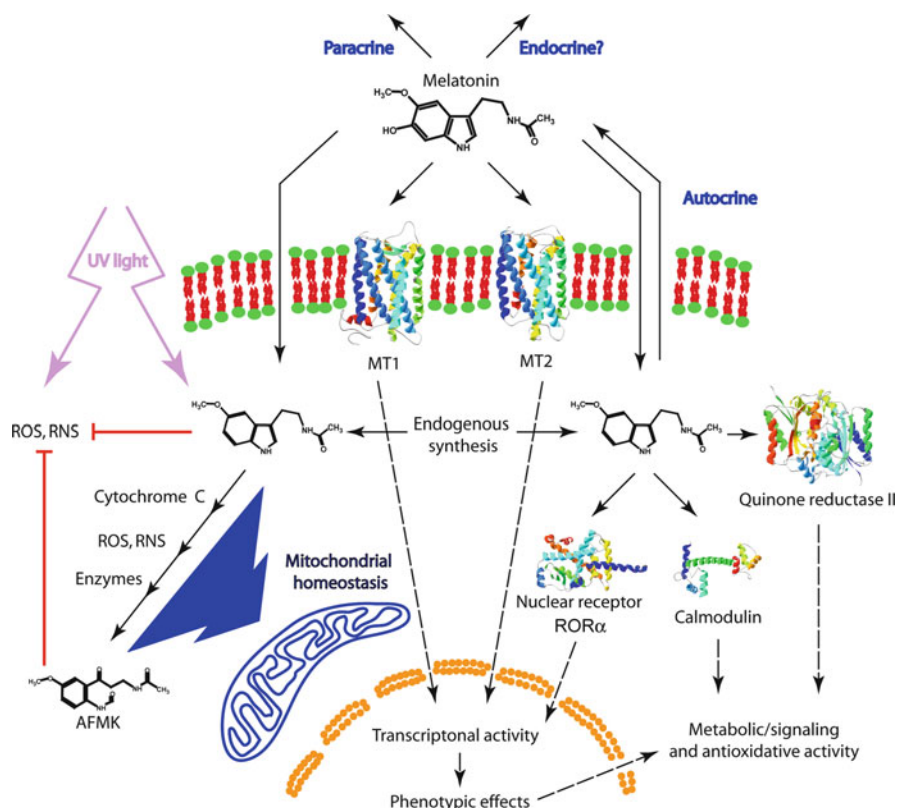


**Fig. 3.1** Pathways of melatonin degradation. The indolic pathway involves 6-hydroxylation of melatonin [I] by CYP1A1, CYP1A2, or CYP1B1 (1) to 6-hydroxymelatonin [II]. Melatonin deacetylase (2) produces 5-methoxytryptamine [III] that is oxidized by monoamine oxidase (3) to form 5-methoxyindoleacetaldehyde [IV], which is converted to 5-methoxyindoleacetic acid [V] by aldehyde dehydrogenase (4) or to 5-methoxytryptophol [VI] by alcohol dehydrogenase (5). In the kynuric pathway, melatonin can be converted non-enzymatically to N1-acetyl-N2-formyl-5-methoxykynuramine (AFMK) [VII] in the process that may include generation of 3-hydroxymelatonin [VIII], 2-hydroxymelatonin [IX], melatonin 2-indolinone [X], 3-hydroxymelatonin 2-indolinone [XI], and melatonin dioxetane [XII] as intermediates. Enzymes or pseudoenzymes (Enz) are involved in melatonin conversion to AFMK. Different pathways can lead to the conversion of AFMK [VII] to AMK [XIII] (6). Reproduced with permission from the publisher (Slominski et al. 2008a)

### 3.3 Biological Activity of Melatonin

At the central level, melatonin functions as chronobiotic regulator (Zeitgeber; circadian pacemaker) regulating photoperiod-dependent reproduction and other biological rhythms, as well as a prominent sleep promoter (reviewed by Pandi-Perumal et al. 2006; Reiter 1991; Watson 1994; Yu and Reiter 1993). It plays a role in reproduction and sexual maturation, energy expenditure, and body mass regulation, acting via central or peripheral receptors (Hardeland et al. 2011; Pandi-Perumal et al. 2006; Reiter 1991; Watson 1994; Yu and Reiter 1993). Melatonin can additionally affect brain and immune, GI, cardiovascular, renal, bone, and endocrine functions. It shows also oncostatic, antiaging, and cell-protective activities (Bartsch et al. 2002; Bubenik 2002; Hardeland et al. 2011; Jung and Ahmad 2006; Luchetti et al. 2010; Pandi-Perumal et al. 2006; Reiter 1991; Watson 1994; Yu and Reiter 1993). Thus, melatonin





**Fig. 3.2** Phenotypic effects of melatonin in the skin. Exogenous or endogenously synthesized melatonin can regulate skin cell phenotype via interaction with melatonin receptors. Non-receptor actions are mediated by interactions with intracellular proteins such as NQO2 or calmodulin. Melatonin and its metabolites can act as direct scavengers of reactive oxygen and nitrogen species (ROS and RNS) and affect mitochondrial functions. Direct effects are shown by *solid lines* and multiple reactions/signaling are shown by *dashed arrow lines*. Reproduced from (Slominski et al. 2008a) with permission from the publisher

has pleiotropic bioactivities acting as a neurotransmitter, hormone, cytokine, and biological response modifier. These functions are mediated by interactions with high-affinity receptors (Dubocovich et al. 2003; Hardeland et al. 2011; Pandi-Perumal et al. 2006; Reiter 1991; Yu and Reiter 1993). The proposed actions of melatonin or its metabolites in the skin are shown in Fig. 3.2.

Melatonin exerts its effects by interacting with specific receptors that are widely distributed throughout the body, and which are differentially expressed in various organs and tissues (Dubocovich et al. 2003, 2010). The cell surface receptor family comprises MT1 (MTNRa) and MT2 (MTNRb) whose coding region genomic structure is similar and constitutes two exons and one intron which show 60% homology at the amino acid level (Dubocovich and Markowska 2005). There are alternatively spliced isoforms of *MT* genes whose functions remain to be defined

(Slominski et al. 2003a, 2005a). It was shown that different genetic variants of *MT2* can affect body glucose homeostasis (Bouatia-Naji et al. 2009; Prokopenko et al. 2009). Both receptors belong to the family of seven-transmembrane receptors coupled to G proteins, i.e., either  $G_i$  or  $G_q/11$ —depending on the receptor type (Dubocovich et al. 2003, 2010). Melatonin activates its receptors at nanomolar or lower concentrations. Binding of melatonin to its receptors affects many cellular signaling pathways. The inhibition of cAMP and cGMP production is followed by the inhibition of CREB, PKA, and c-FOS activities, inhibition of calcium and potassium signaling, modification of protein kinase C activity, stimulation of arachidonic acid release, modification of inositol phosphate turnover, and phosphorylation of the mitogen-activated protein and extracellular signal-regulated kinases 1 and 2 (MEK1 and MEK2) as well as c-Jun N-terminal kinase (JNK) (Capsoni et al. 1994; Dubocovich et al. 2003, 2010).

The oligomerization and dimerization of MT1 and MT2 receptors appear to play a role in the regulation of cell activity (Ayoub et al. 2002; Jockers et al. 2008; Maurice et al. 2010). In addition, melatonin-related receptor (MRR or GPR50), which has high protein sequence homology with MT1 and MT2, may form heterodimers with MT1 (Ayoub et al. 2002; Jockers et al. 2008). Retinoic acid orphan receptor type  $\alpha$  (ROR $\alpha$ ) was proposed to serve as a putative melatonin nuclear receptor (Carlberg et al. 1994; Wiesenberger et al. 1998). However, crystallographic studies have shown that ROR $\alpha$  is in fact a cholesterol sulfate, and not melatonin, receptor (Kallen et al. 2004). Accordingly, others have shown that after binding to MT1 melatonin can indirectly regulate phenotypic activity of ROR $\alpha$  (Hill et al. 2009). It has to be noted that there are at least 4 splicing variants of this nuclear receptor, i.e., *ROR $\alpha$ 1*, *ROR $\alpha$ 2*, *ROR $\alpha$ 3*, *RZR $\alpha$*  (*ROR $\alpha$ 4*) (Becker-Andre et al. 1994; Carlberg et al. 1994; Pozo et al. 2004).

Melatonin exerts also receptor-independent activities. These include broad-spectrum direct antioxidant activity or indirect actions resulting from the activation of anti-oxidative and cytoprotective pathways, a property shared by AFMK and other melatonin metabolites (Hardeland et al. 2009; Reiter et al. 2007; Tan et al. 2001). These properties define melatonin and its metabolites as anti-apoptotic and anti-mutagenic agents (Fischer et al. 2008b; Hardeland et al. 2011; Reiter et al. 2007; Slominski et al. 2008a). Melatonin can also regulate cell metabolism by acting on mitochondria (Hardeland et al. 2011; Semak et al. 2005). Some receptor-independent melatonin actions may be partly mediated via the cytosolic flavoprotein quinone reductase II (NQO2), which is involved in cellular resistance to oxidative stress and detoxification and possesses a melatonin-binding site (previously proposed as a melatonin receptor type 3 (MT3) (Jockers et al. 2008; Nosjean et al. 2000). Melatonin metabolites, including AFMK, and AMK, generated by UVB or oxidative stress can be stronger antioxidants than melatonin itself (Fischer et al. 2006a; Hardeland et al. 2007; Seever and Hardeland 2008). These receptor-independent protective actions of melatonin and its metabolites would require high intracellular levels of the molecules, which can only be met by melatonin in situ production in the relevant tissue, since cellular melatonin uptake is very limited because only 0.1% of extracellular melatonin can enter the cell (Fischer et al. 2006a).

### 3.4 Melatonin Receptors in the Skin

The major compartments of the skin, i.e., the epidermis, dermis, and adnexa, are targets for melatonin regulation (Fischer et al. 2008a; Slominski et al. 2005a, 2008a). More specifically, melatonin was implicated in the regulation of hair growth cycle (Fischer et al. 2008a; Kobayashi et al. 2005; Slominski et al. 2005c), cutaneous pigmentation (Slominski et al. 2004c), as well as skin physiology and pathology (Slominski et al. 2005a) including melanoma (Fischer et al. 2006c; Slominski and Pruski 1993; Yu and Reiter 1993) and vitiligo (Schallreuter et al. 2008a; Slominski et al. 1989). Since those actions have been recently discussed (Dubocovich et al. 2010) the description below will be short. The field of hair growth was the subject of an intensive research in Australia and New Zealand, where experiments on fur-covered animals revealed that melatonin stimulates hair growth. For instance, melatonin-supplemented diet increased the rate of hair growth in springtime (Welch et al. 1990). The results were confirmed in other experimental models (Ibraheem et al. 1994; Nixon et al. 1993). It is likely that fur growth is mediated by melatonin-binding sites/receptors, since these are expressed in rodent skin (Kobayashi et al. 2005; Slominski et al. 1994, 2004b). Clinical studies in women suffering from androgenic alopecia showed a positive effect of melatonin on human hair growth, suggesting it as a potential hair growth regulator in humans (Fischer et al. 2004).

Expression of cell surface MT receptors in the skin varies in different species. To illustrate, skin from the C57BL/6 mouse expresses the *MT2* gene predominantly (Kobayashi et al. 2005) or exclusively (Slominski et al. 2004b), while human skin expresses both MT receptors although a bias toward *MT1* gene expression was observed. The MT1 represents the predominant form of melatonin receptor found in both whole skin and cultured cells (Slominski et al. 2003a, 2005a). As shown by immunocytochemical studies, the expression of MT1 and MT2 proteins in human skin was cell type- and cell compartment-dependent (Slominski et al. 2005a), which suggests that the selectivity of melatonin action could be achieved by spatial compartmentalization and the specificity of signal transduction pathways. The expression of MT receptors was modified by environmental factors in a cell type-specific fashion (Slominski et al. 2003d, 2005a). For example, UVB at 100 mJ/cm<sup>2</sup> induced expression of the *MT1* gene in neonatal epidermal melanocytes, and downregulated it in melanoma cells. *MT1* gene expression was also dependent on the donor, e.g., in some samples of dermal fibroblasts it was not detectable (Slominski et al. 2003d, 2005a). *MT2* gene expression was also upregulated or modified by UVB in epidermal keratinocytes (normal and immortalized), epidermal melanocytes, and dermal fibroblasts. UVB also induced alternatively spliced *MT2b* isoform in epidermal melanocytes, keratinocytes, and dermal fibroblasts (Slominski et al. 2003a). *MT2b* isoforms have two open reading frames (*orf*) encoding the putative proteins. The first *orf*, called MT2b1, would generate a truncated protein of 79 amino acids containing the N-terminal and first transmembrane sequence followed by 8 amino acids of MT2 with the sequence GEHHS added due to a frame shift and the addition of a stop codon. The second *orf*, MT2b2, if translated, would code a protein of 247 amino acid protein, lacking the TM 1-3 domains (Slominski et al. 2003a, 2005c).

Concerning *ROR $\alpha$*  nuclear receptor gene, all of the tested human skin cell types (keratinocyte, melanocyte, and fibroblast lineages) expressed at least one of the three *ROR $\alpha$*  isoforms except for *ROR $\alpha$ 3* gene (Fischer et al. 2006c; Slominski et al. 2005a). Furthermore, UVB downregulated the expression of *ROR $\alpha$*  in HaCaT keratinocytes and upregulated it in normal neonatal melanocytes. *ROR $\alpha$ 1* and *ROR $\alpha$ 2* expression was detected only in dermal fibroblasts and in immortalized melanocytes (*ROR $\alpha$ 2* only) while being undetectable in normal epidermal melanocytes and in keratinocytes (Slominski et al. 2005a). It is likely that *ROR $\alpha$*  gene, due to the alternative splicing (the gene contains 23 exons within a genomic region that spans 732,840 bp), codes other isoforms, as we detected additional DNA fragments of an unexpected length using standard *ROR $\alpha$ 1* and *ROR $\alpha$ 2* primers (Slominski et al. 2005a). The hair cycle-dependent expression of *ROR $\alpha$*  was detected in the mouse skin (Kobayashi et al. 2005). The current challenge is to define whether the RORs still can serve as low-affinity receptors for melatonin taking into consideration that *ROR $\alpha$*  is a specific receptor for sterols (Kallen et al. 2004) and that its role as a receptor for melatonin was questioned (Dai et al. 2001).

### 3.5 Melatonin Protects Against Skin Damage

Serving as an antioxidant and radical scavenger (Tan et al. 2002), melatonin acts as a protecting factor against UVR-induced damage in the skin (reviewed by Fischer et al. 2008b; Slominski et al. 2005a, 2008a). In fact, melatonin is able to prevent sun damage but only when it is administered prior to the UVR exposure and/or is present at the irradiation site (Bangha et al. 1996, 1997; Dreher et al. 1998, 1999). In vitro, melatonin increased cell viability in UV-irradiated fibroblasts (Lee et al. 2003), and decreased apoptosis (Ryoo et al. 2001). Melatonin also protected human leukocytes against UV-induced damage, significantly suppressing ROS formation. It had even stronger radical-scavenging properties than vitamin C and Trolox (Fischer et al. 2002). In human epidermal keratinocytes, melatonin protected against UV-induced reduction of cell viability (Fischer et al. 2006b, 2008c). This effect was found to be receptor-independent (it required high doses of melatonin), and involved anti-apoptotic activities. The interactions between pathways stimulated by UVB and melatonin may be complex since melatonin also attenuated the expression of several genes whose expression was known to be upregulated by UVB (Pisarchik et al. 2004). Thus, melatonin could have a clinically relevant protective action against UVR when used as a sun protective cream component. Although melatonin photostability is a limiting factor, its metabolites 6-hydroxymelatonin and N1-acetyl-N2-formyl-5-methoxykynurenamine can retain significant antioxidant activity (Maharaj et al. 2002). A challenging question is whether protective functions of melatonin depend partly on its regulation of NQO2 function, since NQO2 gene expression is ubiquitous in skin cells (Slominski et al. 2005a).

### 3.6 Conclusions

Melatonin is generated and metabolized in the skin to affect its phenotype as well as to serve as a protective agent against UV radiation. Some of the melatonin effects are mediated through its interaction with melatonin receptors. Other actions result from direct, receptor-independent effects of this free-radical-scavenging molecule as well as metabolic and protective effects induced by melatonin or its metabolites. The pleiotropic activities of the cutaneous melatoninergetic system are mediated by cell-specific intra-, auto-, or paracrine mechanisms, allowing a counteraction or attenuation of both environmental and endogenous stressors leading to the maintenance of skin integrity, and perhaps affecting body's homeostasis (Fig. 3.2). Local melatoninergetic systems could also modify the activities of the cutaneous neuroendocrine network and influence global homeostasis as shown at Figs. 1.1 and 1.2.

## Chapter 4

# Cutaneous Cholinergic System

### 4.1 An Overview

Acetylcholine acts via nicotinic or muscarinic receptors. There are several subtypes of nicotinic receptors that are built of pentamers of at least 17 ( $\alpha 1$ – $\alpha 10$ ,  $\beta 1$ – $\beta 4$ ,  $\gamma$ ,  $\delta$ ,  $\epsilon$ ) subunits (Wu and Lukas 2011). Several subunits of the same type may be present in any given receptor type (e.g.,  $\alpha 1$  and  $\alpha 5$ ). Nicotinic receptors signal by forming ligand-regulated cation channels. There are five subtypes of muscarinic receptors (M1–M5) that act through G-protein-coupled signaling (Graef et al. 2011). Acetylcholine is synthesized by choline acetyltransferase from acetyl coenzyme A and choline. Acetylcholinesterase degrades acetylcholine to acetate and choline. Human keratinocytes synthesize, secrete, and degrade acetylcholine. Choline acetyltransferase is present in all layers of human epidermis, while acetylcholinesterase is present only in basal keratinocytes (Grando et al. 1993). The role of cholinergic system in the skin has been reviewed extensively by Grando and coworkers (2006). The role of acetylcholine-cholinergic receptor system in the skin, which among others regulates the function of eccrine glands, is well known (Fitzpatrick et al. 1993). However, the exclusive role of muscarinic system in sweat glands was challenged by finding nicotinic receptors in myoepithelial and acinar cells of those glands (Kurzen and Schallreuter 2004).

Human keratinocytes express the  $\alpha 3$ -,  $\alpha 5$ -,  $\alpha 7$ -,  $\alpha 10$ -,  $\beta 1$ -,  $\beta 2$ -, and  $\beta 4$ -nicotinic receptor subunits and all types of muscarinic receptors (Grando 1997, 2006; Grando et al. 1995, 1996, 2006). The expression of these receptors changes during the process of keratinocyte differentiation. Basal keratinocytes respond to acetylcholine predominantly via nicotinic receptor  $\alpha 3\beta 2(\beta 4)$  with or without  $\alpha 5$  subunit and the M<sub>2</sub>- and M<sub>3</sub>-muscarinic receptors. Keratinocytes of the prickly layer have more  $\alpha 5$ -containing  $\alpha 3$ -nicotinic receptors, and also express  $\alpha 9$ -nicotinic as well as M<sub>4</sub>- and M<sub>5</sub>-muscarinic receptors.  $\alpha 7$ -nicotinic and M<sub>1</sub>-muscarinic receptors are mainly found on keratinocytes of the granular layer of epidermis (Grando et al. 2006). The  $\alpha 7$ -nicotinic receptor has the most prominent role in keratinocyte differentiation since its deactivation leads to the apoptosis of keratinocytes and the inhibition of

their differentiation. The  $\alpha 3\beta 2$  receptor regulates chemokinesis of leukocytes (Arredondo et al. 2002; Chernyavsky et al. 2004a), while the activation of nicotinic receptors stimulates keratinocyte motility, with  $\alpha 9$  subtype of nicotinic receptor being the most significant in this respect (Nguyen et al. 2000). It was suggested, that, acting simultaneously, nicotinic (primarily  $\alpha 7$ ) and muscarinic (primarily M1) receptors are responsible for directional migration of keratinocytes via the Ras/Raf-1/MEK1/ERK pathway (Chernyavsky et al. 2004a; Grando et al. 2006). Also, the activation of M3-muscarinic receptors favors the expression of migratory integrins and that of M4 promotes sedentary integrins, thereby further solidifying a pivotal role of the cholinergic system in keratinocyte migration and wound re-epithelialization (Chernyavsky et al. 2004b). Muscarinic receptors' activation increased relative amounts of Ki-67, PCNA, and p53 mRNAs as well as PCNA, cyclin D1, p21, and p53 proteins affecting cell cycle (Arredondo et al. 2003).

Acetylcholine's potential role in the pathogenesis of pemphigus can be demonstrated by the fact that cholinergic receptors' activation on keratinocytes altered the expression of desmoglein 1, desmoglein 3, and the phosphorylation status of desmoglein 3 (Nguyen et al. 2003). Cholinergic agonists inhibit the antibody-induced acantholysis and adhesion molecules' phosphorylation in pemphigus vulgaris. M1 ligand binding leads to the activation of both serine/threonine and tyrosine phosphatases, whereas binding of a ligand to the  $\alpha 7$ -nicotinic receptor activates the tyrosine phosphatase and inhibits Src (v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homologue). These processes lead to the dephosphorylation of adhesion molecules and, thus, the inhibition of acantholysis (Chernyavsky et al. 2008).

SLURP family proteins regulate the function of the cholinergic system and their abnormalities are found in one of the palmoplantar keratodermas (Mal de Meleda) and psoriasis (Fischer et al. 2001; Tsuji et al. 2003). Local acetylcholine levels are increased in atopic dermatitis (Wessler et al. 2003).

Human melanocytes express the M<sub>1</sub>–M<sub>5</sub> subtypes of muscarinic receptors and  $\alpha 1$ ,  $\alpha 3$ ,  $\alpha 5$ ,  $\alpha 7$ ,  $\beta 1$ ,  $\beta 2$ ,  $\gamma$ , and  $\delta$  subunits of nicotinic receptors (Buchli et al. 2001). Acetylcholine induces pigmentation via nicotine receptors and inhibits it via M2- and M4-muscarinic receptors (Grando et al. 2006; Kurzen and Schallreuter 2004; Wallstrom et al. 1999). The activity of acetylcholine esterase is decreased in vitiligo (Iyengar 1989). The M4-muscarinic receptor may have a pivotal role in the regulation of murine hair pigmentation (Hasse et al. 2007).

All the elements of the cholinergic system are expressed in Langerhans cells and lymphocytes. Muscarinic M3 receptor expression on lymphocytes is much stronger than other receptor subtypes (Tayebati et al. 2002).

## 4.2 Conclusions

The cholinergic system plays a pivotal role in the regulation of keratinocytes' homeostasis. Their differentiation, motility, adhesion, and cell cycle are modified by acetylcholine. The differential expression of the receptors has been documented

in detail. The  $\alpha 7$ -nicotinic receptor plays a key role in keratinocyte differentiation and  $\alpha 9$ -nicotinic, M3-, and M4-muscarinic receptors in keratinocyte migration. In the melanocytes, the activation of nicotinic receptors induces pigmentation, while the opposite is true for M2- and M4-muscarinic receptors. The cholinergic system is implicated in skin pathologies such as palmoplantar keratoderma (Mal de Meleda type), psoriasis, atopic dermatitis, vitiligo, and pemphigus. In addition, communication between the cutaneous neuroendocrine system and the rest of the body is partly achieved via the cholinergic system (Figs. 1.1 and 1.2).



## Chapter 5

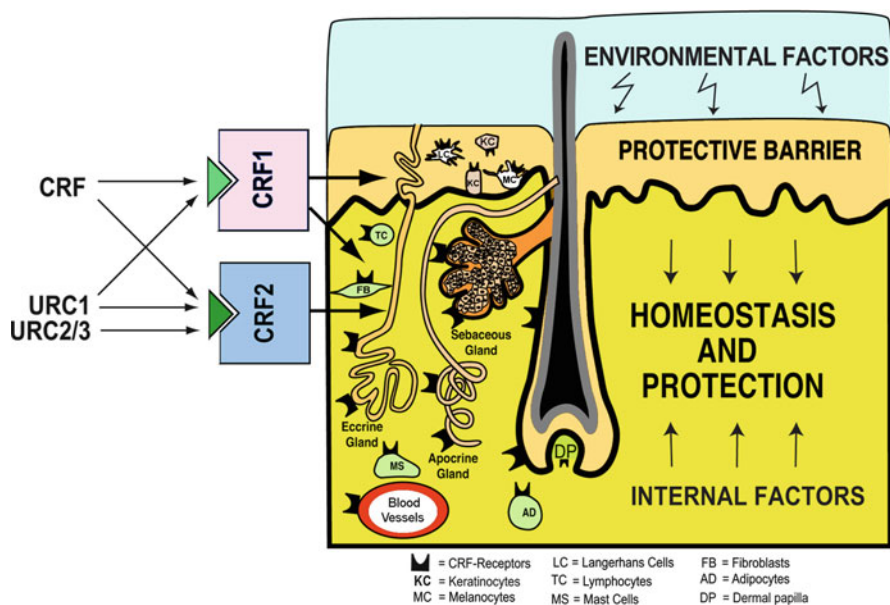
# Corticotropin Signaling System in the Skin

### 5.1 CRF and Urocortins

Corticotropin-Releasing Factor (CRF), a 41 amino acid long hypothalamic neuropeptide discovered by Vale and Rivier (Spiess et al. 1981; Vale et al. 1981), and related urocortin (Urc1-3) are brain neuropeptides that regulate behavioral, autonomic, endocrine, reproductive, metabolic, and immune functions (Grammatopoulos and Chrousos 2002; Hillhouse et al. 2002; Perrin and Vale 1999). In peripheral tissues, they act as local immunomodulators with predominantly proinflammatory actions (Hasse et al. 2007; Slominski 2003b; Slominski et al. 2006c; Theoharides and Cochrane 2004) as well as they directly regulate cardiovascular, gastrointestinal, reproductive, and gestational activities (Hillhouse et al. 2002). These neuropeptides exert their regulatory activities via interaction with CRF receptors, CRF1 and CRF2, which were cloned and initially characterized by Vale's group and others (Grammatopoulos and Chrousos 2002; Hillhouse and Grammatopoulos 2006; Hillhouse et al. 2002; Perrin and Vale 1999; Slominski et al. 2001).

### 5.2 Expression and Functions of CRF and Urocortins in the Skin

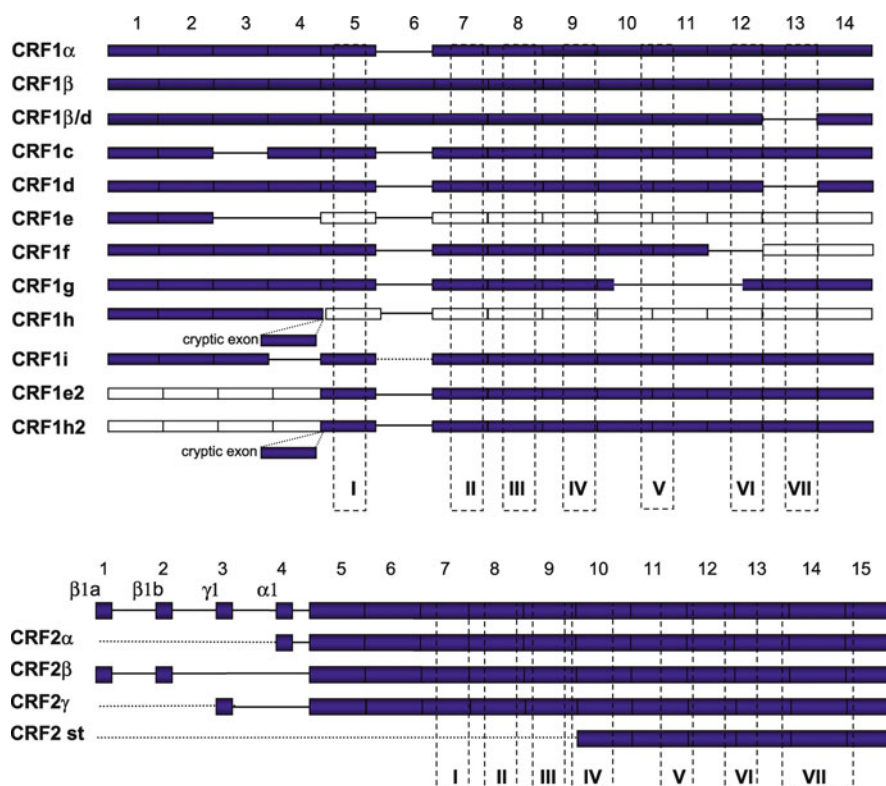
The CRF signaling system regulates human skin homeostasis (Janjetovic et al. 2009; Slominski and Wortsman 2000; Slominski et al. 2000c, 2006c; Zmijewski and Slominski 2009b, 2010) (Fig. 5.1). In fact, we were the first to detect CRF and Urc1 production in the skin (Roloff et al. 1998; Slominski et al. 1996a, 1998b, 1999c, 2000b, c), which was stimulated by UVR and cAMP, and was inhibited by dexamethasone (Slominski et al. 1996a, 1998b). We also identified and characterized the CRF receptors in the skin and defined their functional activity (Slominski et al. 1999c, 2000c, 2001, 2006c), findings confirmed and extended by others. CRF and urocortins can inhibit proliferation of cultured human keratinocytes and melanocytes (Quevedo



**Fig. 5.1** CRF-related signaling in the skin regulates its protective and homeostatic functions. The specificity of the effect is defined either by a local production of molecules (CRF, Urc1, Urc2, or Urc3) or the type of the receptor expressed (CRF1 vs. CRF2). Reprinted from (Slominski 2009b) with permission from the publisher

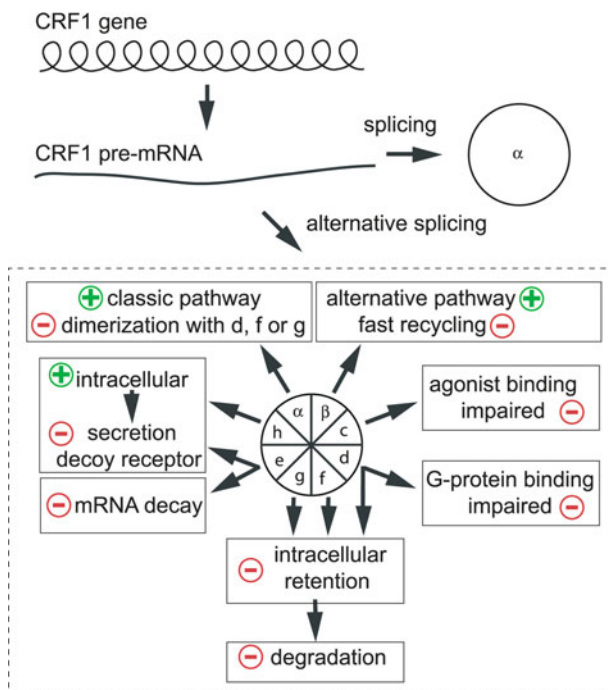
et al. 2001; Slominski et al. 2000b, 2006a; Zbytek et al. 2005). They also stimulate cell differentiation (Zbytek et al. 2005) and modulate the expression of cell surface adhesion molecules and cytokine production by human keratinocytes (Quevedo et al. 2001; Zbytek et al. 2002, 2004). CRF and related peptides stimulate POMC expression and corticosterone and cortisol production (Cirillo and Prime 2011; Hannen et al. 2011; Ito et al. 2005; Rousseau et al. 2007; Skobowiat et al. 2011; Slominski et al. 2005d, e, 2006c; Vukelic et al. 2011; Zbytek et al. 2006b). These phenotypic effects of CRF in the skin are mediated by interaction with CRF1, which is the predominant receptor type expressed in human epidermis, and CRF2, and appear to be secondary to the modulation of intracellular concentrations of cAMP, IP3,  $\text{Ca}^{2+}$ , or NF- $\kappa\text{B}$  activity (Fazal et al. 1998; Slominski et al. 1999c, 2005e, 2006a; Wiesner et al. 2003). Moreover, CRF also stimulates steroidogenic activities in sebocytes which express both CRF1 and CRF2 (Zouboulis et al. 2002). Thus, CRF and Urc exhibit non-endocrine activities like regulation of cell proliferation, differentiation, and immune cell interactions, thereby defining these peptides as a novel type of growth factors/pleiotropic cytokines (Kausar et al. 2006; Slominski et al. 2006a, c; Zbytek and Slominski 2007; Zbytek et al. 2006a).

These pleiotropic activities of CRF and urocortins give the mechanism of regulating CRF signaling in the skin a significant priority (Slominski et al. 1999c, 2006c; Zmijewski and Slominski 2010). CRF1 is expressed in all major cellular lineages of the skin, while CRF1 $\alpha$  prevails in human epidermis, and CRF2 is



**Fig. 5.2** CRF1 and CRF2 receptors and their alternative splicing variants. *Upper panel:* Human CRF1 gene consists of 14 exons and due to alternative splicing at least ten isoforms can be generated with seven found in human skin (Pisarchik and Slominski 2001; Slominski et al. 2006c). Coding exons are shown in blue and noncoding exons due to frame shift followed by in-frame premature stop codon are shown as white squares. *Lower panel:* CRF2 gene contains 15 exons and at least three alternative transcription start codons. Due to alternative splicing at least three main isoforms can be created (CRF1 $\alpha$ ,  $\beta$ ,  $\gamma$ ) and four additional isoforms could be synthesized from full-length mRNA by employing alternative start codons (CRF2 $\alpha$ 1,  $\beta$ 1a,  $\beta$ 1b,  $\gamma$ 1 as shown on the top of the panel). Exon numbers are marked on the top of each panel. Transmembrane segments of the 7TM domains are shown as *squares* (dashed line with number I–VII) (Slominski et al. 2006c; Zmijewski and Slominski 2010)

expressed in all mouse cutaneous compartments (Slominski et al. 2001, 2004a, 2006c). A major challenge in this area is the functional implication of the coupling of different CRF1 isoforms to different signal transduction systems (Figs. 5.2 and 5.3) (Janjetovic et al. 2009; Pisarchik and Slominski 2001, 2004; Slominski et al. 2001, 2004a, 2006a, c; Zmijewski and Slominski 2009a, b, 2010). This differential coupling could provide the mechanistic explanation for observed organ- and cell type-dependent variability in the phenotypic response to CRF as previously suggested (Pisarchik and Slominski 2001, 2004; Slominski et al. 2004a, 2006c; Zmijewski and Slominski 2010).



**Fig. 5.3** Regulation of the CRF signaling by CRF1 isoforms. *CRF1* gene contains 14 exons and only one isoform of the CRF1β receptor (also called pro-CRF1) is coded by all exons. CRF1 transcript is also subjected to alternative splicing resulting in at least eight isoforms. Recent studies showed that the expression and/or co-expression of CRF1 isoforms is responsible for the modulation of CRF1 signaling mediated by main CRF1α or alternative CRF1β isoform. Soluble isoforms (CRF1e and h) were also found to stimulate CRF or modify Urc signaling when co-expressed with CRF1α. “Minus” sign indicates inhibition of CRF signaling on different levels including: fast mRNA decay (CRF1e), dimerization, and subsequent intercellular retention resulting, most probably, in premature receptor degradation (CRF1α with CRF1d, CRFf, or CRFg), decoy receptor mechanism (CRF1h and e when secreted), agonist binding impairment (CRF1c), or G-protein binding inhibition (CRF1d). For details, see (Zmijewski and Slominski 2010). Reproduced with permission from the publisher

### 5.3 Splicing of CRF Receptor Transcripts

Alternative splicing of mRNA is one of the most important mechanisms accountable for genomic variability in higher eukaryotes (Luco et al. 2011). Therefore most human genes, including those coding for G-protein-coupled receptors (GPCRs), are sources for multiple protein isoforms.

CRF receptors are members of class B (secretin family) of G-protein-coupled receptors (GPCRs) and are closely related to calcitonin, growth-hormone-releasing hormone (GHRH), glucagon, glucagon-like peptides, parathyroid hormone (PTH), pituitary adenylate-cyclase-activating peptide (PACAP), and secretin receptors

(Lagerstrom and Schioth 2008). Members of the class B of the GPCR receptor family bind to peptides longer than 27 amino acid residues and are expressed in the majority of endocrine and non-endocrine cells (Hillhouse and Grammatopoulos 2006; Perrin and Vale 1999; Slominski and Wortsman 2000).

In humans, the *CRF1* gene, which contains 14 coding exons, was mapped to chromosome 17 (17q12-q22) (Polymeropoulos et al. 1995). *CRF2* gene has 15 exons (Hillhouse and Grammatopoulos 2006; Slominski et al. 2001) and is located on chromosome 7 (7p14.3). The coding sequences of the two receptors show high degree of homology, although not uniformly distributed along the sequence. A comparison of the protein sequences revealed three distinct regions of homology, corresponding to the structural domains of CRF receptors. Extracellular domain (ECD) of the CRF receptor is responsible for substrate recognition and binding, and this region showed the lowest homology (40%) between *CRF1* and *CRF2*. This feature most probably reflects differential affinity to ligands (CRF and Urc 1-3) (Hillhouse and Grammatopoulos 2006). On the other hand, 7TM domain and intracellular and extracellular loops are highly conserved with homology of around 80%. The most conserved part of the CRF receptor is the third intracellular loop involved in the interaction with G-proteins (Hemley et al. 2007; Hillhouse et al. 2002). Despite a high level of similarity, the pattern of splicing variants of the two known CRF receptor genes (*CRF1* and *CRF2*) seems to be unique for each pre-mRNA (Fig. 5.2).

### 5.3.1 *CRF1* mRNA Splicing Variants

Processing of pre-mRNAs encoding *CRF* receptors may result in an alternative splicing with at least ten variants of *CRF1* mRNA ( $\alpha$ ,  $\beta$ , c, d, e, f, g, h, and i) (Hillhouse and Grammatopoulos 2006; Karteris et al. 2011; Pisarchik and Slominski 2001, 2002; Slominski et al. 2006c, 2007a; Zmijewski and Slominski 2010). Importantly, all of those isoforms except newly discovered *CRF1 $\beta$ /d* and *CRF1i* were found in human skin (Mikhailova et al. 2007; Slominski et al. 2007a; Zmijewski and Slominski 2010, 2011). Interestingly, only one isoform of *CRF1* (*CRF1 $\beta$* ) contains all 14 exons, while the main functional isoform, *CRF1 $\alpha$* , has an exon 6 spliced out. The exon 6 seems to be unique for *CRF1 $\beta$* , although a recent study by Karteris and coworkers (2011) revealed a new isoform, named *CRF1 $\beta$ /d* since it shared the properties of isoform *CRF1 $\beta$*  (exon 6) and *CRF1d* (lack of exon 13). This finding raises a theoretical possibility of the expression of all *CRF1* splicing variants with and without exon 6. Other *CRF1* isoforms might be divided into three groups: soluble receptors (*CRF1e* and *CRFh*), receptors with defects in the ECD (isoform c), and receptors with impaired 7TM domain (*CRF1d*, f, g, and also *CRF1 $\beta$ /d* fits to this group). The detailed exonal organization of *CRF1* isoforms is shown in Fig. 5.2 and was discussed elsewhere (Hillhouse and Grammatopoulos 2006; Slominski et al. 2006c; Zmijewski and Slominski 2010, 2011). It has to be noted that alternative splicing of *CRF1* also results in a frame shift

which introduces premature stop codon to the sequence of CR1e, f, g, and h. Alternative splicing of CRF1 receptor mRNA seems to be a conserved phenomenon in evolution because CRF1 isoforms were also identified in rat (Hillhouse and Grammatopoulos 2006; Slominski et al. 2001), mouse (Pisarchik and Slominski 2001), and hamster (Pisarchik and Slominski 2002). Also, some splice variants are conserved among the members of the family B of GPCRs. For instance, characteristic deletion of exon 13 was found in *CRF1* isoform d and calcitonin receptor (Grammatopoulos et al. 1999; Markovic et al. 2008; Seck et al. 2005; Zmijewski and Slominski 2009b). Theoretically, it is possible that, due to the alternative splicing, also “headless” CRF1 receptor isoforms could be coded by CRF1 pre-mRNA (Slominski et al. 2006c). Their presence was predicted based on the known mRNA sequences of isoforms CRF1e and h (Pisarchik and Slominski 2001; Zmijewski and Slominski 2010). The mRNAs of those isoforms—due to the presence of the alternative code premature stop codon and alternative ATG start cordons—could theoretically allow for the synthesis of “headless” isoforms of CRF1 receptor. Such ECD domain-missing isoforms were identified for the closely related calcitonin receptor (Nag et al. 2007). However, proof for the existence of CRF1 “headless” isoforms remains to be provided.

### 5.3.2 *CRF2 Splicing Variants*

In theory, *CRF2* gene has a capacity for similar number of isoforms as shown for CRF1 (<http://www.ncbi.nlm.nih.gov/IEB/Research/Acembly/>), although only CRF2 $\alpha$ ,  $\beta$ ,  $\gamma$ , and soluble sCRF2 isoforms of *CRF2* were well characterized (Grammatopoulos and Chrousos 2002; Hillhouse et al. 2002). In contrast to *CRF1* gene, the gene of *CRF2* receptor has at least three alternative starting codons coded by alternative exons located on 5'-end of the *CRHR2* gene (see Fig. 5.2). In addition, the presence of the soluble isoform of *CRF2* (sCRFR2 $\alpha$ ) was also reported in mouse brain (Perrin et al. 2003) and headless isoform of *CRF2* was found in stomach (GenBank accession No. E12750; Patent: JP199707289-A).

### 5.3.3 *Expression of CRF1 and CRF2 Isoforms in the Skin*

In human skin, *CRF1* gene is expressed in both epidermal and dermal compartments, whereas *CRF2* is detected predominantly in adnexal structures such as hair follicles (Kausar et al. 2006; Slominski et al. 2004a) or sebaceous glands (Kausar et al. 2006). It seems that this pattern is characteristic for humans because mouse skin expresses both *CRF1* and *CRF2* (Slominski et al. 2004a, 2007a), and both of them take part in the regulation of skin physiology (Kausar et al. 2006; Slominski et al. 2004a, 2006c).

All studied human epidermal and dermal cell lines express the main CRF1 isoform  $\alpha$  and for some cells like adult epidermal keratinocytes and melanocytes, it is the only isoform found under normal conditions (Pisarchik and Slominski 2001; Slominski et al. 2001). Also, melanocytes, keratinocytes, and fibroblasts found in hair follicles express CRF1 $\alpha$  (besides the previously mentioned CRF2) (Kausar et al. 2006; Slominski et al. 2004a, 2006c). However, neonatal epidermal keratinocytes, dermal fibroblasts, and several melanoma cell lines express multiple CRF1 variants (Pisarchik and Slominski 2001; Slominski et al. 2004a).

### **5.3.4 Modulation of the Expression of CRF1 Isoforms and Its Physiological Relevance**

Although the regulation of the alternative splicing of CRF receptor genes' remains unknown, a theoretical model of alternative splicing with a potential involvement of U1 and U2 small nuclear ribonucleoproteins (snRNPs), splicing activators, and Ser/Thr kinases was proposed (Markovic and Grammatopoulos 2009). Here, we will discuss only biological factors which affect *CRF* receptor splicing with relevance to human skin.

The *CRF1* expression pattern and alternative splicing is regulated by diverse physiological and pathological factors, including cell growth conditions or exposure to the ultraviolet irradiation (Zmijewski and Slominski 2010). In human immortalized HaCaT keratinocytes, CRF1 $\alpha$  is the only isoform expressed in confluent culture. However, fast growing (subconfluent) cells express multiple isoforms including  $\alpha$ , c, and e (Zmijewski and Slominski 2009b). In addition, the expression of CRF1 mRNA and protein increases with confluence of HaCaT keratinocyte cultures (Zmijewski and Slominski 2009b). The above phenomena may also explain differences in the CRF1 expression between neonatal and adult epidermal keratinocytes (Slominski et al. 2004a, 2007a).

The pathological conditions can influence the expression of CRF1 receptor as shown in skin biopsies from psoriatic patients (Tagen et al. 2007; Zmijewski and Slominski 2009b). It is worth mentioning that most of the studied melanomas expressed multiple CRF1 isoforms including CRF1 $\alpha$ , except for SKMEL-188 melanoma cells that exclusively expressed CRF1d (Pisarchik and Slominski 2001; Slominski et al. 2004a). This raises a question whether CRF1 splicing is involved in the pathogenesis of skin hyperproliferative (malignant or benign) and inflammatory diseases and whether external and internal stressors can affect skin physiology through context-dependent *CRF1* alternative splicing leading to the differential CRF signaling in this organ (Slominski 2009b; Slominski et al. 2006c; Zmijewski and Slominski 2010).

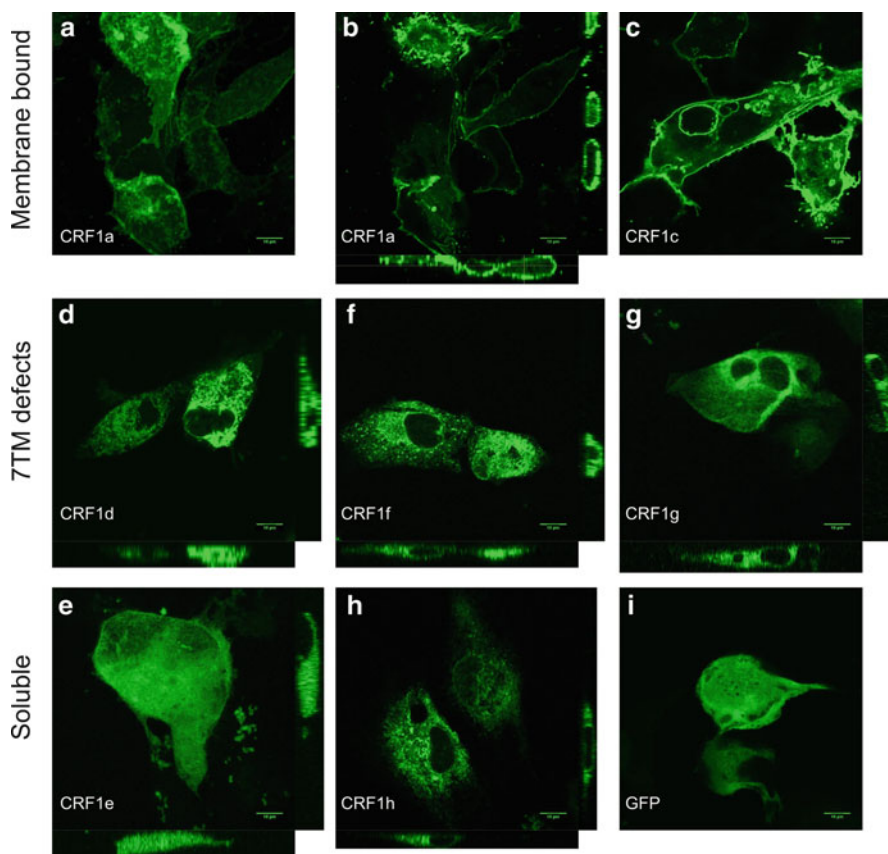
A single-nucleotide polymorphism (SNP) of *CRF1* gene might be an additional factor with high impact on *CRF1* expression and splicing. There is a growing body of evidence that SNPs of *CRF1* are associated with several human pathologies



including hypertension, abusive behavior, and depression (Kamdem et al. 2008; Schmid et al. 2009; Wasserman et al. 2008). Therefore, it is possible that SNPs might influence the *CRF1* gene expression and/or splicing of its pre-mRNA.

Recent studies revealed a potential mechanism and significance of CRF1 isoforms' expression in the epidermal and dermal cell lines and other models (Jin et al. 2007; Karteris et al. 2011; Markovic et al. 2008; Pisarchik and Slominski 2001, 2002; Slominski et al. 2006a, c, 2007a; Sztainberg et al. 2009; Zmijewski et al. 2007; Zmijewski and Slominski 2009b). As stated above, the pattern of CRF1 isoform expression depends on cell type and culture growth condition. These data also correlate with the observed changes in the responsiveness to CRF or urocortin. The current model of the regulation of CRF signaling confirms the central role of CRF1 $\alpha$  and suggests at least modulatory roles for the other CRF1 isoforms in signal transduction (Fig. 5.3) (Zmijewski and Slominski 2010). For example, alternative splicing may decrease the levels of CRF1 $\alpha$  transcripts. Isoform CRF1e is a good candidate. This isoform consists of only first coding exons followed by the premature termination codon in exon 5, resulting in a frame shift due to the removal of exons 3 and 4. This isoform could be subjected to fast decay mechanisms. An additional mechanism of the regulation of CRF1 function by the expression of multiple splicing variants may include heterooligomerization of CRF1 isoforms leading to changes in CRF1 $\alpha$  trafficking to the cell membrane, its localization, and function. Oligomerization of GPCR is a well-known mechanism of the regulation and activation of the GPCR A family. Interestingly, oligomerization of CRF1 isoforms was found in different cell compartments of HaCaT epidermal keratinocytes (Zmijewski and Slominski 2009a) and also in pituitary AtT-20 cells (Zmijewski and Slominski 2009b). These findings confirmed earlier studies which showed dimerization of CRF1 $\alpha$  (Kraetke et al. 2005) and heterodimerization of CRF1 with vasopressin V1b receptor (Young et al. 2007). Thus, the co-expression of CRF1 $\alpha$  with CRF isoforms defective in 7TM domain (d, f, g) may result in the retention of heterodimers inside the cells and thus inhibition of the translocation of the newly synthesized CRF1 $\alpha$  receptors to the cell membrane. Based on structural modeling of CRF1 isoforms, it was concluded that alterations to the sequence caused by alternative splicing should result in the instability of receptors in the cell membrane (Slominski et al. 2006c). Indeed, CRF1d, when overexpressed in HaCaT keratinocytes, localized predominantly to the endoplasmic reticulum, and CRF1f and CRF1g co-localized within Golgi cisterns. Thus, none of the CRF1 isoforms with impaired 7TM domain showed proper membrane localization (Zmijewski and Slominski 2009a, b). The overexpression of those isoforms can influence downstream signaling including cAMP, IP<sub>3</sub> production, and calcium mobilization, followed by altered transcriptional activity (Grammatopoulos and Chrousos 2002; Grammatopoulos et al. 1999; Hillhouse and Grammatopoulos 2006; Markovic et al. 2008; Pisarchik and Slominski 2004; Slominski et al. 2006a, 2007b; Wietfeld et al. 2004; Zmijewski et al. 2007; Zmijewski and Slominski 2009a, b, 2010). The third mechanism of the regulation of CRF signaling may involve direct alterations of the receptor function. CRF1d, f, and g isoforms and their heterodimers with CRF1 $\alpha$ , found predominantly inside the cell, could not





**Fig. 5.4** Localization of the CRF1 isoform tagged with GFP in human adult ARPE-19 cells. Adult retinal pigment epithelium cells (ARPE-19) as alternative to the melanocyte model of pigment-producing cells showed similar intracellular distribution of the CRF1 isoform to that described previously in HaCaT keratinocytes (Zmijewski and Slominski 2009b) and ATT-20 pituitary cells (Zmijewski and Slominski 2009b). Isoforms CRF1 $\alpha$  (Panels **a** and **b**) and CRF1c (Panel **c**) with full-length 7TM are found predominantly within cell membrane. CRF1 isoforms with defects (CRF1d—Panel **d**, CRF1f—Panel **f**, CRF1g—Panel **g**) within 7TM region show intracellular localization. The soluble isoforms (CRF1e—Panel **e**, CRF1h—Panel **h**) are localized predominantly inside the cells. The isoform CRF1e (Panel **e**) is the only isoform found inside the cell nucleus (similarly as GFP alone—Panel **i**). ARPE-19 cells were transfected with constructs carrying CRF1 isoforms fused with GFP (Zmijewski and Slominski 2009c) and images (as Z stacks) were collected with Zeiss LSM 510 laser scanning microscope (Zeiss, Germany). On the bottom and right sides of Panels **b**, **d**, **f**, **g**, **e**, and **h** cross sections (from Z stacks) were shown to emphasize three-dimensional localization of the CRF isoforms. On Panel **a**, Z stack projection (average intensity) of APRE-19 cells overexpressing CRF1 $\alpha$  is shown to emphasize the presence of this isoform on the cell surface. The controls are represented by ARPE cells transfected with GFP alone (Panel **i**)

be activated by extracellular ligands (Pisarchik and Slominski 2004; Slominski et al. 2007b; Zmijewski and Slominski 2009a, 2010). Even if isoforms with impaired 7TM domains (CRF1d, f, and g) would reach proper cell membrane localization, the downstream signaling would be impaired due to improper binding/activation of G-protein (Zmijewski and Slominski 2009a, b, 2010). On the other hand, the isoform CRF1c, despite its proper membrane localization (Slominski et al. 2006c; Zmijewski and Slominski 2009a, b), has a deletion of exon 3 encoding the main part of receptor's ECD, which resulted in impaired ligand binding (Slominski et al. 2006c; Zmijewski and Slominski 2009a). Indeed, CRF1 expressed in COS-1 cells failed to bind antagonist ( $[I^{125}]oCFR$  (Ross et al. 1994). An increase in cAMP production was only observed after stimulation with high concentration of human CRF indicating attenuation of signaling by CRF1c in comparison to CRF $\alpha$  (Karteris et al. 1998).

A fourth type of the CRF signaling modulation is represented by the soluble CRF1h isoform (Slominski et al. 2006c; Zmijewski and Slominski 2011). The intracellular retention and co-localization within ER of CRF1h were shown in cells overexpressing this isoform (Zmijewski and Slominski 2009a, b). In addition, CRF1h was released from the cells to the media and inhibited CRF signaling (Pisarchik and Slominski 2004; Zmijewski and Slominski 2009b). If this process occurs in vivo, it would be consistent with a well-known mechanism of action of soluble receptors (also called decoy receptors) and similar to the function of CRF2 soluble isoform (sCRF2) (Chen et al. 2002). The described models of the CRF signaling regulation via the expression of CRF1 isoforms may be not unique to skin cells but can represent a global mechanism valid for other tissues and cell lineages, because many organs and tissues express multiple CRF1 isoforms (Fig. 5.4).

### 5.3.5 Conclusions

CRF, urocortins, and CRF receptors are widely expressed in human skin. The phenotype of skin cells is affected by endogenously produced CRF and urocortins via a variety of signal transduction systems. Furthermore, expression of multiple CRF1 isoforms may represent additional means of regulating CRF-mediated stress responses at different levels (Fig. 5.2). Different stress signals were found to influence *CRF1* splicing and signaling, but there are also indications that other factors such as small nucleotide polymorphism may influence CRF signaling by modification of the CRF1 pre-mRNA splicing. The recently proposed mechanism of the CRF signaling regulation by its receptor splicing may explain the observed changes in cell responsiveness to CRF and CRF-like ligands (Fig. 5.3).

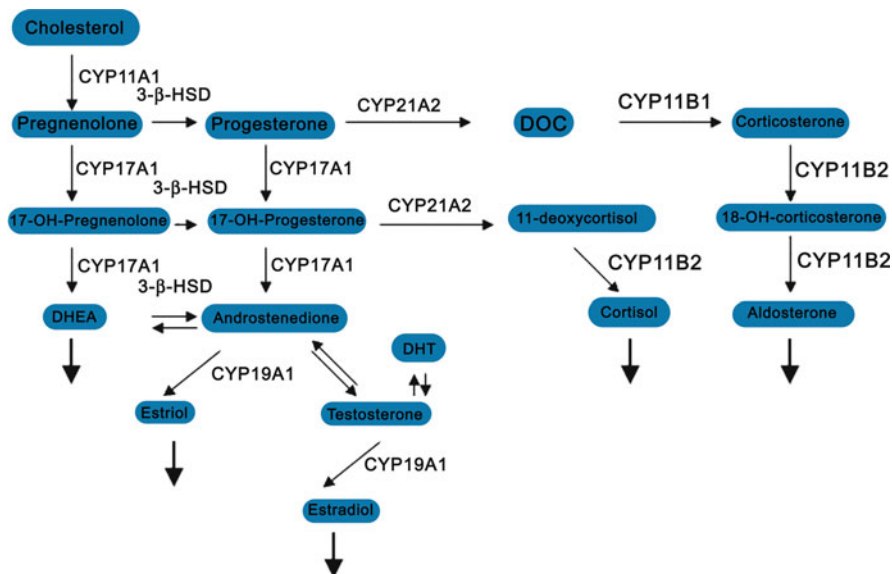
Modulation of *CRF1* splicing may have a direct impact on the differentiation and proliferation of various skin cell types. Alterations of *CRF1* splicing mechanism can ultimately lead to the development or aggravation of symptoms of several skin pathologies including acne, psoriasis, or skin cancer (Fig. 5.1).

## Chapter 6

# Steroidogenesis in the Skin

### 6.1 An Overview

Adrenocortical steroidogenesis is initiated by the interaction of ACTH with melanocortin receptor type 2 (MC2-R) that stimulates secretion and production of cortisol via the activation of steroidogenic enzymes and cholesterol mobilization or transport into mitochondria (Felig and Frohman 2001; Payne and Hales 2004). Prolonged ACTH effects include induction of corresponding enzymes, and the entire process is linked to the ACTH-stimulated cAMP production (Felig and Frohman 2001; Payne and Hales 2004). The biochemical pathway is initiated by the rate-limiting enzyme cytochrome P450<sub>scc</sub> (encoded by the *CYP11A1* gene) that cleaves cholesterol side chain to produce pregnenolone, precursor to all steroids (Payne and Hales 2004; Tuckey 2005). In a classical pathway, pregnenolone is further transformed to corticosterone or cortisol through sequential action of 3 $\beta$ -HSD/isomerase, P450c17, P450c21, and P450c11 $\beta$  enzymes and then released into circulation (Felig and Frohman 2001; Payne and Hales 2004). Specifically, 3 $\beta$ -HSD/isomerase transforms pregnenolone to progesterone, and 17(OH)pregnenolone to 17(OH)progesterone, P450c17 catalyzes 17 $\alpha$ -hydroxylation of pregnenolone or progesterone to 17(OH)pregnenolone or 17(OH)progesterone, and P450c21 hydroxylates progesterone to deoxycorticosterone (DOC) or 17(OH)progesterone to 11-deoxycortisol, while P450c11 $\beta$  hydroxylates DOC to corticosterone and 11-deoxycortisol to cortisol (Felig and Frohman 2001; Payne and Hales 2004; Simard et al. 2005). ACTH also stimulates production of mineralocorticoids and sex hormones via MC2-R and has a trophic effect on the adrenal cortex (Felig and Frohman 2001) (Fig. 6.1). Low-level production of cortisol via the classical pathway was reported in peripheral tissues (Taves et al. 2011) including the gastrointestinal tract, brain (Davies and MacKenzie, 2003; Do Rego et al., 2009), immune cells (Costa et al. 2009; Vacchio et al. 1994), and also in colon cancer (Sidler et al. 2011). In addition, cortisol levels are also regulated by 11 $\beta$ -HSD1 (Draper and Stewart 2005), which at high NADPH/NADP<sup>+</sup> ratios transforms cortisone to cortisol (Draper and Stewart 2005; Tomlinson et al. 2004). Cortisol can also be



**Fig. 6.1** A scheme of the steroidogenic pathway

transformed to cortisone by  $\text{NADP}^+$ -dependent enzyme  $11\beta\text{-HSD2}$  that acts exclusively as a dehydrogenase (Tomlinson et al. 2004). The expression of both  $11\beta\text{-HSD1}$  and  $11\beta\text{-HSD2}$  enzymes was shown in placenta, kidney, liver, fibroblasts, and adipocytes (Bujalska et al. 1997, 2002; Ricketts et al. 1998; Tiganescu et al. 2011).

## 6.2 Cutaneous Corticosteroidogenic System

We were the first to demonstrate that human skin expresses genes encoding enzymes involved in the sequential metabolism of cholesterol to pregnenolone and to corticosteroids including cytochromes *P450<sub>scc</sub>*, *P450<sub>c17</sub>*, and *P450<sub>c21</sub>*, and the *MC2-R* (receptor for ACTH) genes (Slominski et al. 1996d). These findings were later complemented by the demonstration of these enzymes' functional activity in epidermal and dermal skin cells (Rogoff et al. 2001; Slominski et al. 1999b, 2000a, 2002d, 2004a) and that cutaneous steroidogenesis begins from cholesterol (Slominski et al. 2004d, 2007a). Rapid metabolism of progesterone and deoxycorticosterone (DOC) was shown in rodent skin and cultured human normal and malignant skin (Slominski et al. 1999b, 2000a, 2002d). Thus, production of corticosterone and DOC-like steroid species was shown in rat skin (Slominski et al. 2000a). Also cultured malignant melanocytes showed progressive transformation of progesterone to DOC, 18-hydroxy-DOC, and corticosterone, but not to aldosterone (Slominski et al. 1999a). Cortisol and corticosterone production

was further documented in normal epidermal melanocytes (Slominski et al. 2005e) and dermal fibroblasts (Slominski et al. 2005d, 2006b), with final evidence on cortisol synthesis by human skin cells provided by liquid chromatography-mass spectrometry (LC/MS) analysis (Slominski et al. 2005e, 2006b). In agreement, cortisol production was demonstrated in human hair follicles (Ito et al. 2005; Sharpley et al. 2009), and cortisol production by epidermal and dermal skin cells was later confirmed by other authors (Cirillo and Prime 2011; Hannen et al. 2011; Vukelic et al. 2011). Interestingly, cutaneous cortisol production was mediated by both CYP11B1 and 11 $\beta$ -HSD1 activities (Cirillo and Prime 2011; Hannen et al. 2011; Slominski et al. 2007a; Tiganescu et al. 2011; Vukelic et al. 2011). While some authors found cortisol production by early passages of human epidermal keratinocytes (Cirillo and Prime 2011; Hannen et al. 2011; Vukelic et al. 2011), we (Slominski et al. 2005e) and others (Milewich et al. 1986) did not detect cortisol in late passages of epidermal keratinocytes or melanocytes. In agreement with the last finding, in HaCaT keratinocytes, progesterone and DOC were metabolized rapidly to steroid products different from corticosterone, aldosterone, and cortisol (Slominski et al. 2002d). These discrepancies may be due to the contamination of primary cultures of keratinocytes by other cell types (e.g., melanocytes) or differences in culture conditions. Cortisol production by skin cells was regulated by ACTH and factors raising cAMP level (Slominski et al. 2005e), IL-1, as well as by wound response (Vukelic et al. 2011) and high energy ultraviolet radiation (Skobowiat et al. 2011).

### 6.3 Production of Sex Hormones in the Skin

The skin is an important organ transforming dehydroepiandrosterone (DHEA) and DHEA-sulfate (DHEA-S) or androstenedione, which predominantly originate from systemic circulation, to active sex hormones (Labrie et al. 2003; Ohnemus et al. 2006; Zouboulis 2004; Zouboulis and Degitz 2004; Zouboulis et al. 2007) (Fig. 6.1). In addition, in the skin local steroidogenic system (Ito et al. 2005; Slominski and Wortsman 2000; Slominski et al. 2002d, 2004d, 2005d, e, 2008b; Taves et al. 2011) produces 17(OH)pregnenolone and 17(OH)progesterone that are further metabolized to DHEA with their following metabolism to androgens and estrogens (Fig. 6.1), or other steroidal products (Slominski et al. 2002d, 2009a, c).

DHEA of systemic or local origin is transformed by 3 $\beta$ -HSD into 4-androstenedione, and 5-androstene-3 $\beta$ ,17 $\beta$ -diol into testosterone, while 17 $\beta$ -HSD converts DHEA into 5-androstene-3 $\beta$ ,17 $\beta$ -diol, 4-androstenedione into testosterone, and androstenedione into DHT (Labrie et al. 2000, 2003; Milewich et al. 1991; Simard et al. 1993; Zouboulis et al. 2008; Zouboulis and Degitz 2004). Cutaneous testosterone is also converted into dihydrotestosterone (DHT) by the action of a 5 $\alpha$ -reductase (reviewed by Zouboulis et al. 2008). Skin and subcutaneous adipose tissue is also an important site of estrogen production, in particular after menopause (Labrie et al. 2003; Ohnemus et al. 2006; Zouboulis et al. 2007).

Furthermore, it is an important site of estrogen and androgen activation (Fig. 6.1) (Labrie et al. 2000; Ohnemus et al. 2006; Zouboulis et al. 2007; Zouboulis and Degitz 2004). The locally produced sex hormones modify skin phenotype and function via interactions with the corresponding androgen and estrogen receptors (Labrie et al. 2003; Ohnemus et al. 2006; Randall et al. 1993; Slominski and Wortsman 2000; Zouboulis 2004; Zouboulis et al. 2007; Zouboulis and Degitz 2004).

## 6.4 Conclusions

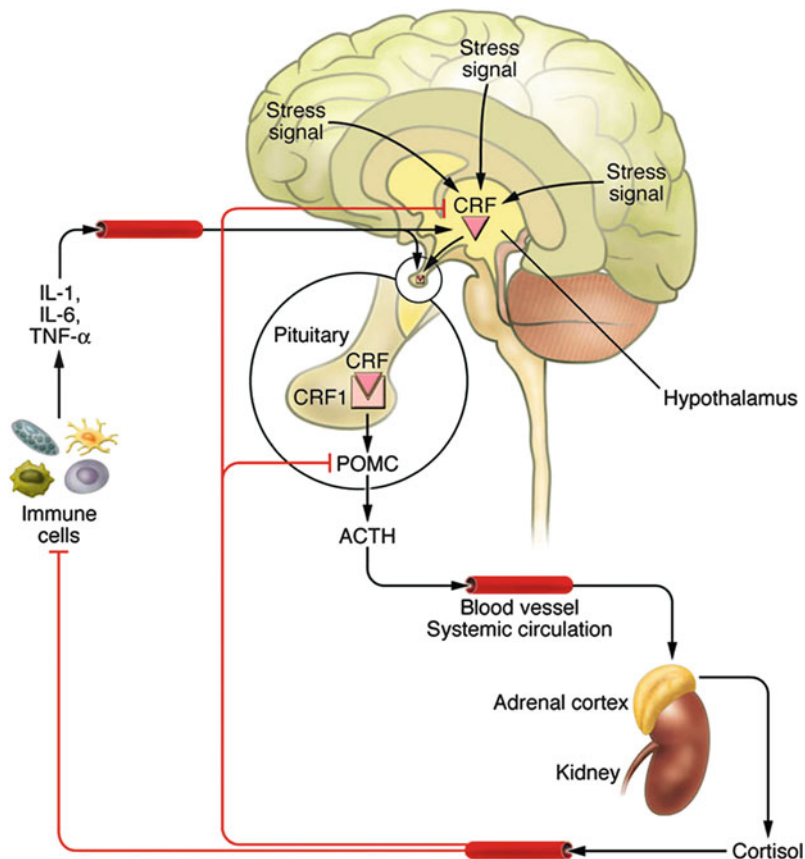
Mammalian skin is an extra-adrenal site of mineralo/glucocorticoid synthesis, which can be regulated by endogenous and environmental factors (Slominski et al. 2007a, 2008a). Furthermore, the skin is an important site for estrogen and androgen production, activation, or metabolism (Labrie et al. 2003; Ohnemus et al. 2006; Zouboulis et al. 2007). Interestingly, skin production of steroids seems to be cell type dependent and subjected to the regulation by external factors such as ultraviolet radiation. These steroids act in intra-, auto-, or paracrine fashions to regulate local homeostasis. Moreover, skin and its subcutaneous tissue constitute an important source of estrogens and androgens in females, especially after menopause.

## Chapter 7

# Equivalent of Hypothalamo–Pituitary–Adrenal Axis in the Skin

### 7.1 Systemic HPA Axis

The work of Hans Selye was fundamental in defining the hypothalamic–pituitary–adrenal (HPA) axis as the body's important coordinator of responses to systemic stress (Selye 1936; Seyle 1976). The HPA functional structure has been completed by determining that hypothalamic corticotropin-releasing factor (CRF) acts as the regulator of the production of ACTH and  $\beta$ -endorphin in the anterior pituitary (Spiess et al. 1981; Vale et al. 1981). The HPA pathway (Fig. 7.1) is triggered by various stress factors which activate production of CRF in the paraventricular nucleus (PVN) (Chrousos 1995; Chrousos and Gold 1992; Owens and Nemeroff 1991). In pituitary CRF binds to CRF type 1 receptors (CRF1) (Aguilera et al. 2001; Hillhouse and Grammatopoulos 2006; Perrin and Vale 1999) increasing production and secretion of the proopiomelanocortin (POMC)-derived peptides, i.e., ACTH, MSH, and  $\beta$ -endorphin (Hillhouse and Grammatopoulos 2006; Pritchard and White 2007; Smith and Funder 1988). The arginine vasopressin (AVP) produced by the PVN can also act synergistically with CRF in activating the HPA axis (Chrousos 1995; Itoi et al. 2004). In the adrenal cortex ACTH, by interacting with the MC2 receptors (MC2-R), stimulates production and secretion of cortisol in humans or corticosterone in rodents. These corticosteroids counteract the effects of stressors by mobilization of energy reserves, buffering tissue damages, and suppressing immune system. Moreover, corticosteroids via feedback mechanisms inhibit the HPA axis through the suppression of CRF and POMC production. The HPA axis is also controlled by cytokines, tissue modifiers, and growth factors, which can be either produced in the brain or by peripheral tissues including cells of the immune system. Thus, there are various ways of controlling stress responses at the level of hypothalamus or pituitary that bypass the central brain coordinating centers (c.f. in Besedovsky and Rey 2007; Blalock and Smith 2007; Chesnokova and Melmed 2002) (Fig. 7.1).

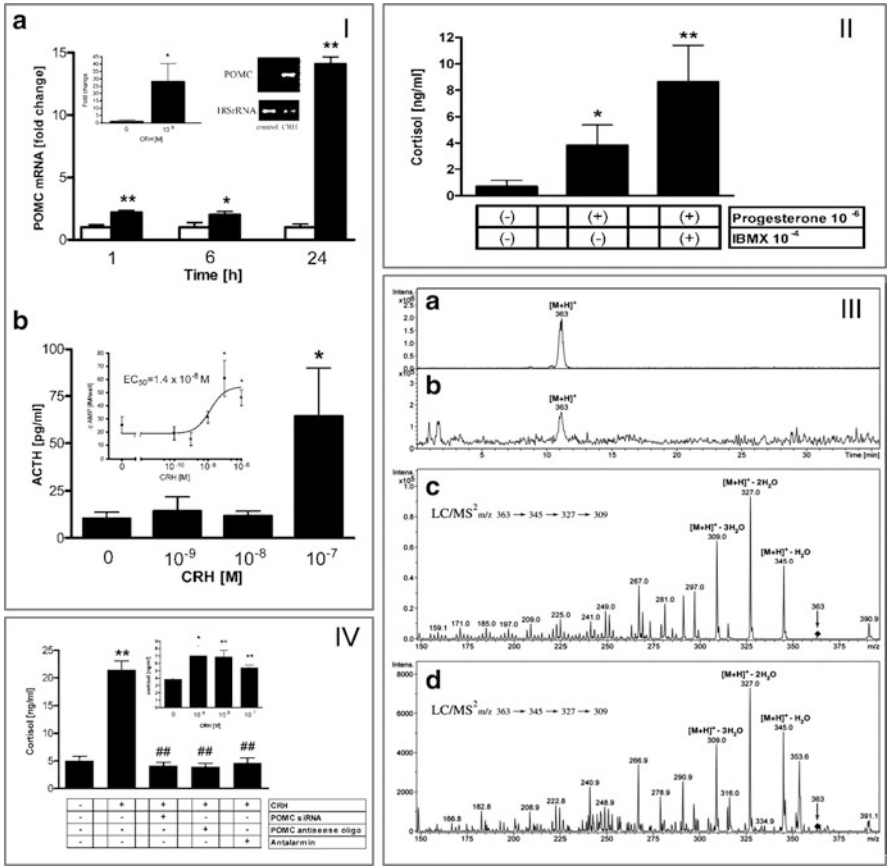


**Fig. 7.1** Organization of the systemic HPA axis. Modified from Fig. 1 published in Slominski (2007)

## 7.2 HPA Axis Homologue Is Expressed in the Skin

More than a decade ago, we proposed that skin expresses a homologue of the HPA axis to regulate local stress responses (Slominski et al. 1996a). This concept was based on finding all molecular elements of the HPA axis in the mammalian skin, i. e., CRF, CRF1, POMC, ACTH, MC2-R, glucocorticoid receptors, and genes coding steroidogenic enzymes (Slominski 1991; Slominski et al. 1992, 1993b, c, 1995, 1996c, 2000d). Over the last 15 years, our and other laboratories provided definitive evidence that skin expresses CRF and the POMC-derived  $\beta$ -endorphin, ACTH and  $\alpha$ -MSH, the corresponding CRF1, melanocortin (MC) and opiate receptors, along with the key enzymes of corticosteroid synthesis, which results in the cutaneous production of corticosterone and cortisol (Arck et al. 2006; Rogoff et al. 2001; Slominski et al. 2000c, 2004d, 2006b, 2007a; Slominski and Wortsman 2000; Tobin 2006; Tobin and Kauser 2005b). Furthermore, we presented data

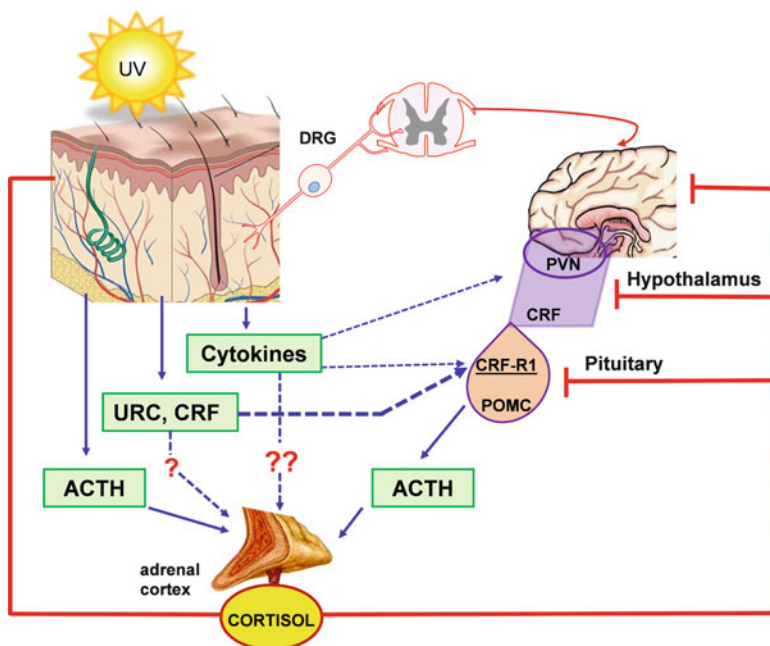




**Fig. 7.2** HPA algorithm is expressed in normal human melanocytes. (a) CRH stimulates POMC gene expression and ACTH production with attendant stimulation of cAMP. (b) Cortisol production is enhanced by the addition of progesterone and/or IBMX (inhibitor of phosphodiesterase). (c) Cortisol is identified by LC/MS<sup>2</sup> in melanocytes (standard = a, c; conditioned media from melanocytes = b, d). (d) CRH stimulates cortisol production that is dependent on POMC expression and CRF1 signaling. Reproduced from Slominski et al. (2005e) with permission from the American Physiological Society

indicating that CRF can stimulate cortisol production in skin cells via POMC cleavage products (Fig. 7.2) (Ito et al. 2005; Slominski et al. 2005e, 2006b, 2007a). These studies provided accumulating evidence that the cutaneous stress response system follows the functional hierarchy of the central HPA with its direct local phenotypic consequences and systemic implications (Slominski et al. 2007a, 2008b; Slominski 2009b) (Fig. 7.3).

Importantly, CRF, POMC, and corresponding receptors were co-expressed in cultured keratinocytes, melanocytes, or dermal fibroblasts (Slominski et al. 2000c, 2006b) as well as their co-expression was demonstrated in vivo in the skin by in situ hybridization or immunocytochemistry (Funasaka et al. 1999; Ito et al. 2005;



**Fig. 7.3** Skin stress response system can activate the central HPA with its direct homeostatic, metabolic, and phenotypic effects. We hypothesized that global responses (at body's level) to UVR may be initiated in the skin and involve a simultaneous activation of sensory receptors and local production of humoral messages (Slominski 2005; Slominski and Wortsman 2000; Slominski et al. 2008b). These signals are either delivered by ascending nerve routes to the brain or by circulation to hypothalamus to activate CRF production in the PVN; CRF then would enter hypophyseal portal circulation with a subsequent activation of CRF1 in the pituitary. Skin humoral signals can also enter directly the pituitary from the circulation. The final outcome for these signaling processes results in the stimulation of ACTH release and of POMC activity. Alternatively, although less likely, the cutaneous factors can bypass the HPA and enter adrenal gland directly from the circulation. The net effect of all these processes is the release of cortisol/corticosterone and induction of steroidogenesis with subsequent metabolic and homeostatic effects

Kauser et al. 2006; Kono et al. 2001; Rogoff et al. 2001; Slominski et al. 2000c, 2006b). Finally, the expression of the executive arm of the HPA, i.e., production of cortisol and corticosterone, has been clearly demonstrated in cultured epidermal keratinocytes and melanocytes as well as in dermal fibroblasts (Hannen et al. 2011; Slominski et al. 2005d, e, 2006b; Vukelic et al. 2011). Moreover, production of cortisol and corticosterone was also shown in human hair follicles maintained in vitro (Ito et al. 2005; Sharpley et al. 2009), and in rodent skin. In fact, skin contains an entire biochemical apparatus necessary to transform cholesterol to cortisol and corticosterone (Slominski et al. 2007a) including the capability to produce pregnenolone (Slominski et al. 2004d; Thiboutot et al. 2003) and its further sequential transformation to progesterone, deoxycorticosterone (DOC), 18-hydroxy-DOC, cortisol, and corticosterone (Dumont et al. 1992; Ito et al. 2005; Rogoff et al. 2001; Slominski and Wortsman 2000; Slominski et al. 1999b, 2000a,

2005d, e, 2006b). The cutaneous expression of the above HPA axis elements is nonrandom, but is organized into functional, cell type-specific regulatory loops with a structural hierarchy similar to the central HPA (Slominski et al. 2007a) (Fig. 7.3). Specifically, exogenous CRF interacted with CRF1 on cultured human epidermal melanocytes and dermal fibroblasts stimulating cAMP production with subsequent increases of POMC gene expression and production of ACTH (Slominski et al. 2005d, e, 2006b). Similarly, CRF stimulated POMC production in immortalized normal and malignant melanocytes expressing CRF1 receptor (Slominski et al. 2006b; Zbytek et al. 2006a), and CRF stimulated POMC and  $\alpha$ MSH production by epidermal and follicular melanocytes (Rousseau et al. 2007). Most importantly, normal human melanocytes responded to CRF, ACTH, and factors raising intracellular cAMP with an increased production of cortisol and corticosterone, which was dependent on functional CRF1 receptor, since CRF1 receptor's antagonists abolished the effect, and on POMC expression, because silencing of the POMC gene abolished this effect (Fig. 7.2) (Slominski et al. 2005e). Thus, melanocytes, cells of neural crest origin, not only produce CRF, but also respond to it following an algorithm of the central HPA axis, adjusted to the local environment. While fibroblasts also responded to CRF and ACTH with enhanced production of corticosterone, cortisol levels were not regulated by axis, since cortisol was produced constitutively (Slominski et al. 2006b), thus indicating a partial departure from the classical algorithm of the HPA regulation.

The HPA components were also demonstrated in organ-cultured human scalp skin (Ito et al. 2005) where exogenous CRF increased POMC expression with sequential increases in ACTH and cortisol production. These were consistent with in situ co-localization of CRF, CRF1, and POMC in the skin and hair follicles (Kono et al. 2001; Rogoff et al. 2001; Slominski et al. 2000c). Exogenous cortisol inhibited CRF, MC2-R, and ACTH expression by interacting with glucocorticoid receptors (Ito et al. 2005) which was in agreement with earlier studies which showed inhibition of the POMC and CRF expression by dexamethasone (a synthetic glucocorticoid) in mouse skin (Ermak and Slominski 1997) and cultured human skin cells (Slominski et al. 1998b). Thus, mammalian skin expresses a fully functional HPA axis equivalent which encompasses local CRF, ACTH, and cortisol/corticosterone synthesis, and secretion with a negative feedback regulation of CRF and POMC expression mediated by glucocorticoid receptors' activation (Slominski et al. 2007a). The stimulatory role of ACTH on cortisol production by human epidermal cells has been confirmed recently (Cirillo and Prime 2011; Vukelic et al. 2011).

It is likely that the CRF-driven patterns of steroidogenic responses can be differential depending on cell subpopulation, their tissue localization, and microenvironment (Fig. 5.1) (Slominski et al. 2007a). We have proposed a crucial role of paracrine communications in the skin where keratinocytes, melanocytes, fibroblasts, immune cells, and nerve endings can serve as signal initiators (CRF, ACTH, cortisol, or corticosterone) and recipients (binding to corresponding receptors) (Slominski 2005). The latter function would include an active and compartment-specific intercellular cross-talk and exchange of intermediates of the steroidogenic pathway.

### 7.3 Regulation of the Cutaneous HPA Axis

To serve a role of a coordinator and executor of local responses to stress, cutaneous homologue of the HPA should be activated by specific physical, chemical, and biological skin stressors in organized fashion encompassing local production of CRF and POMC-derived peptides which interact with their respective receptors (Fig. 7.3). Indeed, an exposure of skin or skin cells to UVR stimulated in a time- and dose-dependent manner expression of CRF and POMC genes which was followed by the production and release of CRF,  $\beta$ -endorphin, and ACTH peptides, expression of CRF1, PC1, MC2, MC1, CYP11A1, and CYP11B1, and production of cortisol (Chakraborty et al. 1999; Pisarchik and Slominski 2001; Skobowiat et al. 2011; Slominski et al. 1996b; Zbytek et al. 2006b). The stimulatory effects were predominantly seen after exposure to UVB or UVC with only limited responses to UVA (Skobowiat et al. 2011).

In epidermal melanocytes, the UVB-induced stimulation of the *CRH* promoter was suppressed by both the inhibitors of protein kinase A (PKA) and a plasmid overexpressing dominant mutant CREB (Zbytek et al. 2006b). Accordingly, UVB stimulated CREB phosphorylation, the binding of phosphorylated CREB to CRE sites in the CRF promoter, and the activity of the reporter gene construct driven by consensus CRE sites, while the mutation in the CRE site of the *CRF* promoter rendered the corresponding reporter gene construct less responsive to UVB (Zbytek et al. 2006b). In addition, pharmacological inactivation of CRF1 by selective inhibitors abrogated the UVB-stimulated induction of POMC (Zbytek et al. 2006b). These results indicate that UVR induces CRF1 signaling by stimulating the PKA pathway with the subsequent stimulation of POMC production, which imitates HPA's organizational structure. Our most recent results also show that the ability to activate or modify the "cutaneous HPA" elements is dependent on highly energetic UV wavelengths (UVC and UVB) implying a dependence on their noxious activity (Skobowiat et al. 2011).

### 7.4 Functional Activity of the Cutaneous HPA

#### 7.4.1 Local Effects

All of the HPA elements (CRF and/or POMC signaling systems and steroidogenic activities), separately or in concert, can have profound phenotypic effects in the skin (Figs. 5.1 and 7.3) (Slominski et al. 2000c, 2006c, 2007a) and may affect systemic body responses via neuroendocrine and hormonal signal transmission (Slominski 2005; Slominski et al. 2008b). In the skin, these interactions can follow the classical pathway  $\text{CRF} \rightarrow \text{CRF1} \rightarrow \text{POMC} \rightarrow \text{ACTH} \rightarrow \text{corticosterone/cortisol}$ . However, the context-dependent departures from this central algorithm such as  $\text{CRF} \rightarrow \text{CRF1}$ ,  $\text{CRF} \rightarrow \text{CRF1} \rightarrow \text{POMC}$ ,  $\text{POMC} \rightarrow \text{ACTH} + \text{MSH} + \beta\text{-END}$

and POMC  $\rightarrow$  corticosterone/cortisol may take place. The important local phenotypic outcomes of the entire axis or its departures are protective measures against environmental stressors (physical, biological, and chemical insults). This results from fine-tuning and selective regulation of skin pigmentation, barrier function, adaptive and innate immunity, and adnexal structures' activity. The elements of the cutaneous HPA can also counteract skin pathology, such as inflammatory and autoimmune disorders as well as hyperproliferative and dysplastic processes, in order to protect and restore skin homeostasis. In this context, non-endocrine activities of CRF and related urocortins make CRF signaling in the skin an important regulatory system. Similarly, a central role is assigned to the insult-regulated POMC expression and its context-dependent processing in the skin because the chemical nature of the final peptides defines the phenotypic effect (Slominski et al. 2000c, 2004d, 2007a). Finally, local steroidogenic activity would protect skin homeostasis and counteract pathologic processes but also, on the other hand, terminate protective responses to prevent their potential dyshomeostatic effects (Slominski 2009b; Slominski et al. 2007a, 2008b).

### 7.4.2 Systemic Effects

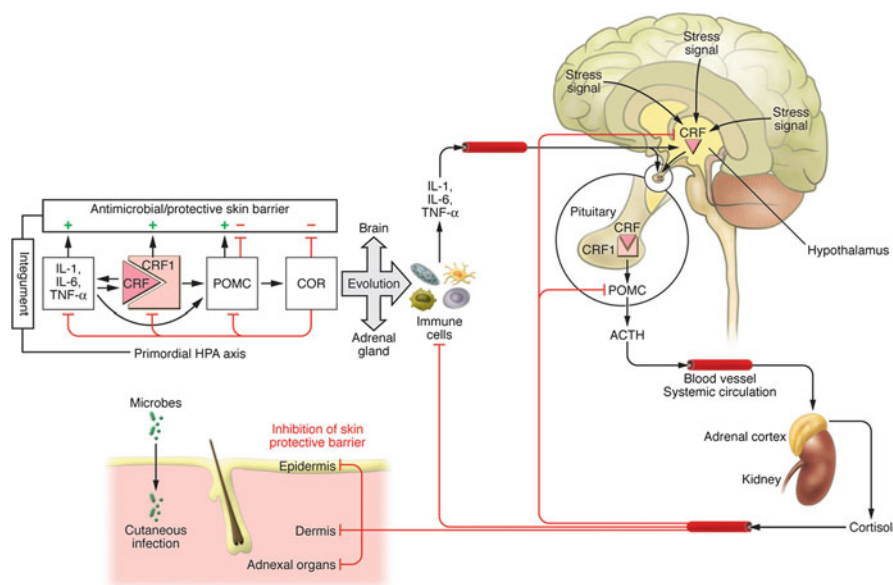
The possibility of the communication between local and systemic HPA axes as well as the differential activation of skin-derived axis elements poses an interesting research question (Slominski 2005; Slominski et al. 2008b) due to the apparent evolutionary conservation of a similar organization at both central and peripheral levels (Slominski 2007). Although all environmental factors noxious to the skin could participate in this communication (Slominski and Wortsman 2000), the role of solar radiation in illustrating the above-mentioned communication and regulation of body homeostasis is the most instructive (Fig. 7.3) (Slominski et al. 2008b). Namely, UVR regulation of systemic homeostasis via HPA could include stimulation of CRF synthesis in the hypothalamus along neutrally transmitted signals from the skin. An increased hypothalamic CRF release would automatically lead to the activation of the existing HPA with cortisol/corticosterone serving as final effector. The HPA axis could be entered at the level of hypothalamus, pituitary, or adrenal gland by skin-derived humoral messages including, respectively, cytokines (action on hypothalamus and/or pituitary), CRF/Urc 1 (action on the pituitary and, possibly, adrenal gland), or ACTH (action on the adrenal gland). This type of regulation would represent a fundamental paradigm shift in neuroendocrinology and photobiology with profound implications for clinical medicine (see above and below).

These concepts are underscored by the observation that humans and horses exposed to sunlight led to increased serum levels of  $\alpha$ -MSH and ACTH (Holtzmann et al. 1982, 1983), while experimental whole body exposure to UVB increased  $\beta$ -LPH and  $\beta$ -endorphin serum levels (Belon 1985; Levins et al. 1983). This model may provide mechanistic explanation for a well-known phenomenon of systemic immunosuppressive action of UVB (Kripke 1994) or serve as an alternative

explanation for the reported cases of the attenuation of multiple sclerosis in some patients after exposure to UVR, a phenomenon so far linked to the increased production of vitamin D (Becklund et al. 2010). Thus, our model of the UVR-mediated activation of central HPA axis may serve as a rational background for a phototherapy of systemic autoimmune disorders or other pathologies. Lastly, our model may provide mechanistic explanation of the recently described phenomenon of “UVR addiction” (Kourosh et al. 2010; Nolan et al. 2009) caused by cutaneous  $\beta$ -endorphin production.

## 7.5 Common Origin of the Central and Peripheral HPA

Evolutionary conservation of a similar HPA-like organization at central and peripheral levels has been documented (Arck et al. 2006; Slominski 2005; Slominski and Wortsman 2000; Slominski et al. 2000c, 2001, 2004d, 2006c, 2007a, 2008b). The common ectodermal origin of brain and epidermis raises the fundamental question of whether the peripheral CRF signaling system is an evolutionary duplicate of its central homologue or whether the brain itself has adopted the preexisting peripheral CRF response system during evolution of the central nervous and endocrine systems. Since it had been shown that cytokines and growth factors can modify CRF and POMC-related functions in pituitary and brain (Slominski and Wortsman 2000; Slominski et al. 2006c) and that CRF can also act as a growth factor/cytokine [a function that develops at the periphery (Kausar et al. 2006; Slominski et al. 2006a, 2006c; Zbytek and Slominski 2007)], we proposed a new hypothesis on the integumental origin of the HPA axis (Slominski 2007). We suggest that the primordial HPA (Fig. 7.4) had first developed in the integument to regulate its defensive activity against the hostile environment and pathogens. Key elements of this system include CRF-related peptide(s) that coordinate innate immune activity and skin barrier formation via CRF1 (an integrating receptor) and thus, both directly and indirectly, affect the expression of the proinflammatory cytokines such as IL-1 and TNF- $\alpha$ . The feedback inhibitory loop begins with CRF1-activated POMC-derived production/secretion and culminates with the production/secretion of corticosterone/cortisol that “shuts off” HPA axis activity and inhibits skin barrier activity. The intermediate signaling molecules (POMC peptides) can both weaken the skin protective barrier by their immunosuppressive action and strengthen it by stimulating melanogenesis as well as direct antimicrobial effects. Thus, the protective barrier functions could be regulated and fine-tuned by the primordial HPA, because of the close association of all of its elements. During evolution, the main algorithm CRF > CRF1 > POMC > ACTH > corticosterone/cortisol may have been adapted and perfected by the central neuroendocrine system to form the HPA axis which has separated anatomically and functionally from the immune system and the skin (Slominski 2007; Slominski et al. 2007a, 2008b). In this context, the retained cutaneous HPA may serve as an evolutionary record of the primary system (Slominski 2007) and, paradoxically, the systemic stress response can weaken the



**Fig. 7.4** Proposed evolution of the stress response system. Reproduced with permission from the publisher (Slominski 2007)

cutaneous antimicrobial defenses as a result of cortisol/corticosterone release from the adrenal glands (Aberg et al. 2007) (Fig. 7.4). It is also possible that both systems had derived from a common precursor and evolved in parallel, maintaining neuro-immune-endocrine communication during this process that has helped to preserve this fine organization of stress responses at the systemic and local levels.

## 7.6 Conclusions

Many important elements of local and systemic responses to environmental stressors (biological, chemical, and physical with predominant role of UVR) originate in the skin, and they involve multiple pathways encompassing activation of different components of the cutaneous and systemic HPA. Hence, signals generated by the integrated actions of CRF, POMC peptides, and cortisol/corticosterone may counteract the local effects of the environment. Furthermore, depending on the type of the stressor and its intensity, the skin can activate systemic HPA either via neural transmission by afferent nerve fibers to the brain or by skin-derived factors which may activate pituitary gland or directly act on the adrenal cortex (Fig. 7.3).



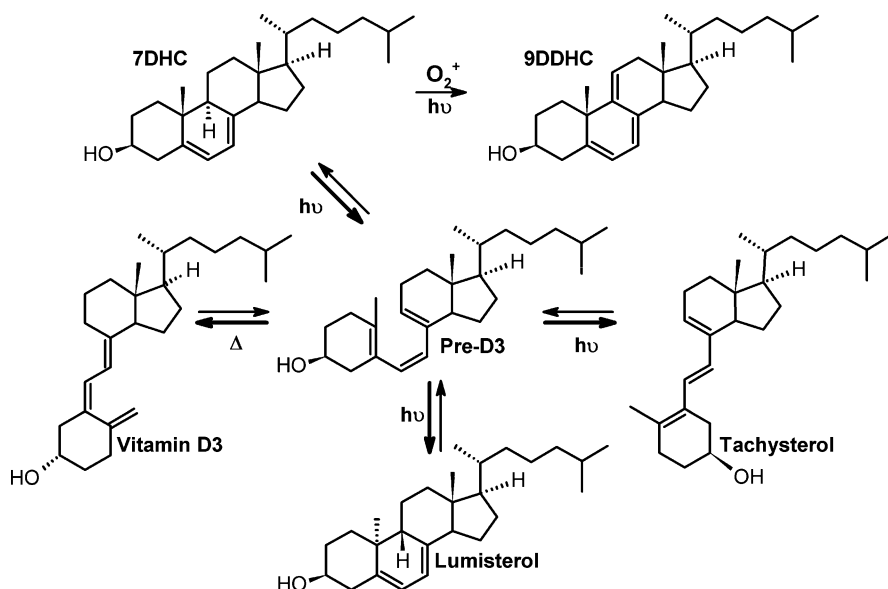
## Chapter 8

# Cutaneous Secosteroidal System

### 8.1 7- $\Delta$ Steroids

Few years ago, it was shown that the P450<sub>scc</sub> system cleaves the side chain of 7-dehydrocholesterol (7DHC, provitamin D<sub>3</sub>) to produce 7-dehydropregnenolone (7DHP) (Guryev et al. 2003; Slominski et al. 2004d). 7DHP is a substrate for a novel metabolic pathway for the synthesis of steroids with two double bonds in 5 and 7 positions, collectively called steroidal 5,7-dienes (Fig. 8.1), which, hypothetically, may be produced in the skin since cutaneous CYP11A1 (P450<sub>scc</sub>) expression was confirmed (Slominski et al. 2004d). The role of this new pathway was supported *ex vivo* by demonstrating the efficient metabolic transformation of 7-dehydrocholesterol to 7DHP by adrenal glands and by mitochondria isolated from rat skin (Slominski et al. 2009c). HPLC separations, UV spectra, and mass spectrometry identified 7DHP, 22-hydroxy-7DHC, and 20,22-dihydroxy-7DHC as the major products with additional minor products defined as 17-hydroxy-7DHP and 7-dehydroprogesterone (Slominski et al. 2009c). These findings defined a novel steroidogenic pathway: 7DHC  $\rightarrow$  22(OH) 7DHC  $\rightarrow$  20,22(OH)<sub>2</sub>7DHC  $\rightarrow$  7DHP, with potential further metabolism of 7DHP mediated by 3 $\beta$ -HSD or CYP17 along the  $\Delta^4$  and  $\Delta^5$  steroidogenic pathways, with the production of 7-dehydroprogesterone and 17(OH)7DHP as intermediates (Slominski et al. 2009c). The existence of this synthetic pathway is documented by the accumulation of pregna- and androsta-5,7-dienes and their hydroxylated derivatives in the Smith–Lemli–Opitz syndrome (SLOS), characterized by 7DHC  $\Delta$ -reductase deficiency, an enzyme responsible for the conversion of 7DHC to cholesterol (Marcos et al. 2004; Shackleton et al. 1999, 2002; Tint et al. 1994). Most recently, we have found that human placenta *ex utero* can transform 7DHC to 7DHP and further to 7-dehydroprogesterone (Slominski et al. submitted for publication).



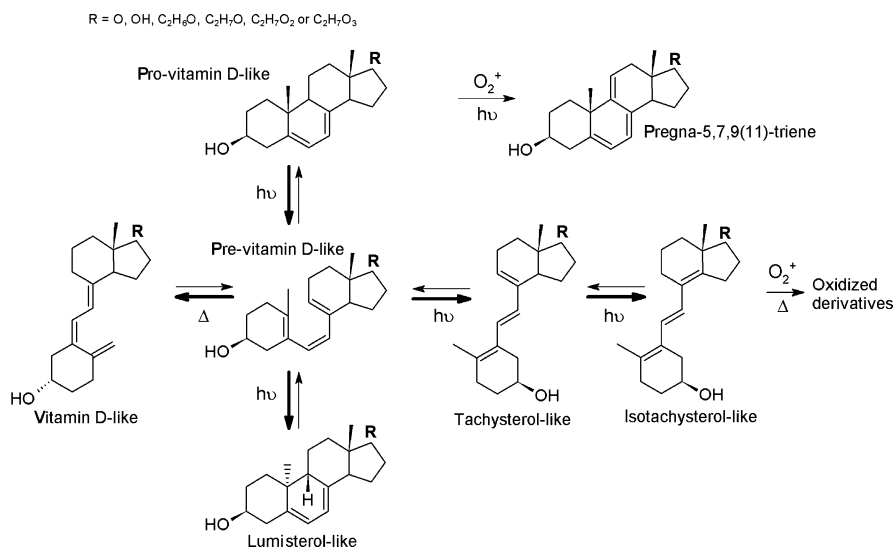


**Fig. 8.1** UVB-induced production and transformation of 7-dehydrocholesterol. Reprinted from Slominski (2009a) with permission from the publisher

## 8.2 Secosteroidogenesis

The UVB-driven isomerization of 7DHC to vitamin D<sub>3</sub> [(3b,5Z,7E)-9,10-secocholesta-5,7,10(19)-trien-1a,3b,25-triol] is one of the most fundamental chemical reactions in vertebrates (Holick 2003; Holick and Clark 1978; Holick et al. 1995). This conversion is initiated by photolysis of the unsaturated B ring on absorption of UVB solar energy of 290–320 nm wavelength producing a pre-D<sub>3</sub> intermediate, followed by its slow isomerization to three main products: vitamin D<sub>3</sub>, tachysterol<sub>3</sub>, and lumisterol<sub>3</sub>. The rate of isomerization depends on the dose of absorbed UVB energy, wavelength, temperature, and the presence of biological membranes (Fig. 8.1) (Buchli et al. 2001; Holick et al. 1995; Tian and Holick 1999). In response to high doses of UVB, another pathway of vitamin D<sub>3</sub> degradation/isomerization yields 5,6-transvitamin D<sub>3</sub>, as well as suprasterols I and II were also demonstrated in the skin (Buchli et al. 2001). In biological systems provitamin D<sub>3</sub> (7DHC) can also be transformed to 5,7,9(11)-trienes in a chemical process that involves an interplay between singlet oxygen and photosensitizers (Albro et al. 1994; Chignell et al. 2006; De Fabiani et al. 1996; Feng et al. 2006; Valencia and Kochevar 2006).

The 5,7-dienal steroids described above as well as P450scc-derived hydroxyl derivatives of 7DHC undergo UVB-induced transformation to androsta-calciferols (aD), pregna-calciferols (pD), and novel hydroxyderivatives of vitamin D<sub>3</sub> (Kim and Lee 2010; Li et al. 2010; Slominski et al. 2004d, 2009a, 2009c; van Beek et al. 2008; Zmijewski et al. 2011) (Fig. 8.2). Furthermore, P450scc



**Fig. 8.2** UVB-induced transformation of prena- or androsta-steroidal 5,7-dienes.  $h\nu$  - photon's energy in reaction, Reprinted from Slominski (2009a) with permission from the publisher

also metabolizes vitamin  $D_3$  to  $20(OH)D_3$ ,  $20,23(OH)_2D_3$ ,  $22(OH)D_3$ ,  $20,22(OH)_2D_3$ ,  $17,20,23(OH)_3D_3$ , and several other  $D_3$ -hydroxyproducts. Novel P450<sub>scc</sub>-derived secosteroids show anti-proliferatory and pro-differentiation activities in a cell type-restricted fashion that depends on the length of the side chain (Janjetovic et al. 2009, 2010; Li et al. 2010; Nguyen et al. 2009; Slominski et al. 2009a, c, 2010, 2011d; Zbytek et al. 2008; Zmijewski et al. 2009a, 2010). These compounds are anti-tumorigenic, can stimulate keratinocyte differentiation, and inhibit NF- $\kappa$ B, acting by binding to vitamin D receptor (VDR) as its partial receptor agonists. They are as potent as  $1,25(OH)_2D_3$ , however, unlike  $1,25(OH)_2D_3$ , only weakly stimulate CYP24 expression (Janjetovic et al. 2009, 2010; Slominski et al. 2009a; Zbytek et al. 2008). Importantly,  $20(OH)D_3$  and  $20,23(OH)_2D_3$  did not affect calcium homeostasis at concentrations as high as 3–4  $\mu$ g/kg (Slominski et al. 2010, 2011b). Thus, we discovered novel metabolic pathways initiated by the enzymatic action of cytochrome P450<sub>scc</sub> (CYP11A1) that produces biologically active novel secosteroids or their precursors, the systemic and local significance of which, including their occurrence in the skin, remains to be defined.

### 8.3 Vitamin D Activity in the Skin: An Overview

The role of skin in the physiology and pathology of vitamin  $D_3$  and its derivatives was a subject of extensive reviews; therefore, below we present only short overview and refer the reader to more extensive descriptions (Bikle 2011a, b, c; Denzer et al. 2011;

Field and Newton-Bishop 2011; Holick 2003, 2008; Lehmann et al. 2004; Pinczewski and Slominski 2010; Reichrath 2007).

The epidermal keratinocytes not only are the site of the photochemical transformation of 7-dehydrocholesterol to vitamin D<sub>3</sub> but also possess the entire enzymatic machinery capable of activating and inactivating vitamin D<sub>3</sub> and its derivatives. Vitamin D<sub>3</sub> is activated by sequential hydroxylation in position 25 by CYP27A1 and in position 1 $\alpha$  by CYP27B1 to form calcitriol, i.e., 1,25(OH)<sub>2</sub>D<sub>3</sub> [(1 $\alpha$ ,3 $\beta$ ,5Z,7E)-9,10-secocholesta-5,7,10(19)-trien-1,3,25-triol]. 1,25(OH)<sub>2</sub>D<sub>3</sub> is inactivated by the action of CYP24 yielding 1,24,25(OH)<sub>3</sub>D<sub>3</sub>. The phenotypic effects of vitamin D<sub>3</sub> and some of its derivatives are mediated by its interaction with VDR which is expressed in all skin cells including keratinocytes, melanocytes, fibroblasts, and other resident cells of the skin. VDR belongs to the family of nuclear receptors and has ligand-activated pleiotropic activities including inhibition of cell proliferation, stimulation of cell differentiation, and modulation of immune functions of skin resident and immigrant cells, to name the most important. Vitamin D is also involved in the regulation of skin barrier function, modulation of skin stress responses, regulation of hair follicle cycling, and suppression of tumorigenesis. To exert those pleiotropic effects, VDR, after dimerization with RXR (retinoic acid receptor X) and translocation to the cell nucleus, interacts with the D receptor-interacting protein (DRIP), the steroid receptor co-activator (SRC) family proteins (with SRC2 and 3 expressed in keratinocytes),  $\beta$ -catenin, and the inhibitor hairless protein (Hr) (Bikle 2011b).

The gradient of calcium level in the epidermis, defined by its low content in the basal layer and the highest level in the corneal layer, determines the expression of several genes required for differentiation of the epidermis and proper epidermal barrier formation. The calcium-stimulated keratinocyte differentiation requires the activity of several proteins including calcium receptor (CaR), phospholipase PLC- $\gamma$  1, and SRC kinases (Bikle 2011a). Calcitriol, the active form of vitamin D<sub>3</sub>, was shown to stimulate the expression of CaR, which subsequently sensitizes keratinocytes to calcium. Moreover, calcitriol is involved in the regulation of synthesis, processing, and trafficking of glycosylceramides, which are important for the regulation of epidermal permeability barrier. In human keratinocytes calcitriol induces the expression of loricrin, filagrin, and phospholipase PCL- $\gamma$ 1 and, synergistically with Ca<sup>+2</sup>, increases the expression of involucrin and transglutaminase K (enzyme required for the cornification of keratinocytes). Anti-proliferative effects of calcitriol are also mediated by inhibiting c-myc, cyclin D1, p21, and p27 (Bikle 2011b).

The pleiotropic effects of vitamin D<sub>3</sub> on skin physiology cannot be solely explained by the differential expression of VDR, which is the highest in the basal layer of the epidermis. As previously mentioned, the activity of VDR is modulated by its interactions with DRIP coactivator complex and SRC proteins. Moreover, it was also shown that in undifferentiated keratinocytes VDR preferentially binds to DRIP coactivator complex in contrast to the more differentiated epidermal cell layers, where VDR interacts with SRC proteins (Bikle 2010).

Vitamin D is also involved in the regulation of skin immune responses by stimulating the expression of I $\kappa$ B $\alpha$ , a NF- $\kappa$ B inhibitor, which leads to the retention of NF- $\kappa$ B in the cytoplasm and its subsequent degradation (Janjetovic et al. 2009, 2010; Lu et al. 2004; Reichrath 2007). Thus, vitamin D and its derivatives inactivate the major transcription factor responsible for the transcription and release of many inflammatory mediators. On the other hand, vitamin D can stimulate innate immune responses, including production of antimicrobial peptides (Gombart et al. 2005; Reichrath 2007).

Basal and squamous cell carcinomas are the most frequent types of human cancer. Because vitamin D and its derivatives inhibit proliferation and stimulate differentiation of keratinocytes, its possible use in the treatment of skin cancer and hyperproliferative diseases of the skin, including psoriasis, has been suggested (reviewed by Bikle 2011a; c; Holick 2008; Lehmann et al. 2004). It is likely that hedgehog or  $\beta$ -catenin pathways can serve as targets of the anticarcinogenic activity of vitamin D in the skin (Bikle 2011c; Tang et al. 2011; Teichert et al. 2011). Furthermore, calcitriol and other derivatives of vitamin D show anti-melanotic activities, and defective expression of VDR correlates with poor prognosis of melanoma (reviewed by Berwick et al. 2005; Brozyna et al. 2011; Field and Newton-Bishop 2011; Janjetovic et al. 2011; Pinczewski and Slominski 2010; Slominski et al. 2011b).

## 8.4 Conclusions

We discovered a novel pathway initiated by the enzymatic action of CYP11A1 on 7DHC (provitamin D3), vitamin D3 (Slominski et al. 2004d, 2009c), and plant-derived, i.e., provitamin D2 (Slominski et al. 2005g, 2006d; Tuckey et al. 2011). The CYP11A1 metabolites of 7DHC, and hydroxyproducts of 7DHP generated by enzymes of classical steroidogenic pathway to produce 5,7-dienal steroids, can be converted by ultraviolet B radiation (UVB) to novel vitamin D hydroxyderivatives, androsta-calciferols, and pregna-calciferols, which are biologically active in keratinocytes and melanoma cells (Janjetovic et al. 2010; Slominski et al. 2005f, 2010, 2011b; Zbytek et al. 2008; Zmijewski et al. 2011). In addition, lumisterol- and tachysterol-like compounds are produced in the skin by the action of UVB. The biological activity of the vitamin D-derived compounds is defined by their chemical structure and cell lineage. Since P450 $_{scc}$  is expressed in the skin, we propose that novel secosteroidal pathways may affect skin biology with potent therapeutical implications. Moreover, body's global homeostasis can potentially be affected by skin-derived secosteroids as well as by CYP11A1-mediated hydroxylation of hydroxyvitamin D in organs or tissues which exhibit high activity of this enzyme.

The associations of vitamin D levels with various pathological states, including osteoporosis, rickets, coronary heart disease, and carcinogenesis, have recently

become the subject of intensive clinical investigations. It is hoped that the results of these studies will provide sufficient data to define the requirements for vitamin D dietary supplementation. Also, the newly described low calcemic vitamin D derivatives, such as 20-OH D<sub>3</sub>, may serve in the future for anticancer therapy.

## Chapter 9

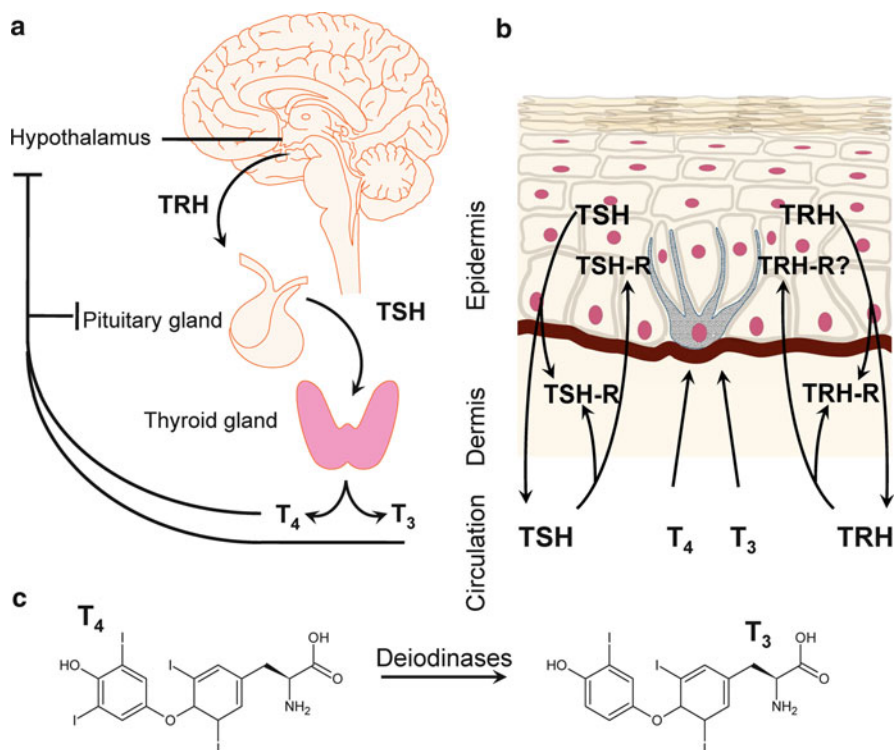
# Equivalent of Hypothalamic–Pituitary–Thyroid Axis

### 9.1 Overview

The hypothalamic–pituitary–thyroid (HPT) axis represents the central regulatory mechanism of cellular metabolism including protein, carbohydrate, and lipid catabolism. The HPT axis is also involved in cell differentiation and proliferation as well as morphogenesis. Tight control of thyroid hormone activity is orchestrated by hypothalamic synthesis of the thyroid-releasing hormone (TRH), which activates its receptors (TRH-R1) in the anterior pituitary, followed by thyroid-stimulating hormone (TSH) secretion. TSH stimulates the production of 3,5,3'-triiodothyronine (T3) and thyroxine (T4) by thyrocytes. Elevated serum levels of thyroid hormones inhibit TRH and TSH synthesis by hypothalamus and pituitary, respectively (Fig. 9.1). Although T4 is the major thyroid hormone found in circulation, T3 is a fully active hormone since deiodinases (type 2 and 3) are expressed in a variety of tissues, including skin, and facilitate the conversion of T4 to T3 (Fig. 9.1). Skin is a nonclassical target for TSH, TRH, and thyroid hormones.

### 9.2 HPT in the Skin

The expression of the molecular elements of the HPT axis (genes for TSH-R1, TSH, TRH, TRH-R1, deiodinases 2 and 3, thyroglobulin, and sodium iodide transporter) in human skin and functional TSH receptors in keratinocytes and malignant melanocytes was first shown by Slominski et al. (2002e). Follow-up studies demonstrated TRH and TSH receptors' expression in various skin cell types including keratinocytes, melanocytes, fibroblasts, and hair follicles (Bodo et al. 2010; Cianfarani et al. 2010; Gaspar et al. 2010; Pattwell et al. 2010; van Beek et al. 2008), which gave ground to the concept of a cutaneous HPT axis that would show similarities to and differences with the central HPT axis (Slominski et al. 2002e). The presence of TSH and TRH receptors explains the phenotypic changes in



**Fig. 9.1** Structure and functions of an equivalent of the hypothalamic–pituitary–thyroid axis (HPT) in the skin

epidermal and dermal cells treated with TRH and TSH (Bodo et al. 2010; Cianfarani et al. 2010; Gaspar et al. 2010; Pattwell et al. 2010; Slominski et al. 2002e; van Beek et al. 2008). The interaction of T<sub>3</sub> with its receptors (TR $\alpha$  and TR $\beta$ ) affects epidermal differentiation and enhances its responsiveness to growth factors (Billoni et al. 2000; Slominski and Wortsman 2000). These effects of T<sub>3</sub> are particularly important for the function of sebaceous, eccrine, and apocrine glands, growth of hair follicles, and synthesis of proteo- and glycosaminoglycans by dermal fibroblasts. For instance, both types of T<sub>3</sub> receptors can regulate keratinocytes' proliferation, differentiation, and immune activity (Contreras-Jurado et al. 2011). Results of the latter study suggest that thyroid hormones acting via their receptors can inhibit skin inflammation, most likely by inactivating specific transcription factors: AP-1, NF- $\kappa$ B, and STAT3 (Contreras-Jurado et al. 2011). T<sub>4</sub> stimulates the proliferation of hair follicle keratinocytes and T<sub>3</sub> inhibits their apoptosis (van Beek et al. 2008). Moreover, thyroid hormone receptors might suppress invasiveness and metastatic ability of skin tumors as shown in mouse knockout models (Martinez-Iglesias et al. 2009). Thyroid hormones may also affect hair follicle stem cells, since T<sub>3</sub> and T<sub>4</sub> were found to induce differentiation and apoptosis, and inhibit clonal growth of hair follicle epithelial stem cells (Tiede et al. 2010). The activity of

deiodinases enables cutaneous conversion of T4 into T3, what plays a role in the regulation of the proliferation of keratinocytes and dermal fibroblasts in vitro and in vivo (Huang et al. 2011; Safer et al. 2009).

The effects of thyroid hormones in the skin are well pronounced in case of thyroid gland pathology including autoimmune thyroid disease (Cianfarani et al. 2010; Slominski and Wortsman 2000). Dermal manifestations of hyperthyroidism include erythema, palmoplantar hyperhidrosis, acropathy, and infiltrative dermopathy. Moreover, Graves' disease also may be associated with generalized itching, chronic urticaria, presence of alopecia areata, and vitiligo (Doshi et al. 2008; Ingordo et al. 2011; Kasumagic-Halilovic et al. 2011). Skin manifestations of thyroid disorders are in part correlated with elevated serum levels of thyroglobulin (Tg), thyroperoxidase (TPO), and thyroid-stimulating hormone receptor (TSH-R) antibodies (Cianfarani et al. 2010; Slominski and Wortsman 2000). Conversely, in hypothyroidism the skin is cool and dry with pasty appearance; hair are commonly dry, coarse, and brittle with up to a 50% probability of diffuse or partial alopecia development (reviewed by Slominski and Wortsman 2000). Several studies also underlined the significance of TSH and TRH in the human skin and their influence on hair physiology (Bodo et al. 2010; Cianfarani et al. 2010; Gaspar et al. 2010). The physiological activity of TRH was demonstrated in human hair follicles' organ cultures expressing active TRH-R1 receptor. TRH stimulated hair shaft formation, prevented apoptosis, increased proliferation of hair matrix keratinocytes, and prolonged the anagen phase of hair growth cycle (Gaspar et al. 2010; van Beek et al. 2008). TRH can also stimulate hair follicle pigmentation (Gaspar et al. 2011), probably by direct activation of melanocortin type 1 receptor (MC1-R), confirming our previous hypothesis (Slominski et al. 2002e, 2005b). It was shown that TSH acting via TSH-R1 receptor increased cAMP production by human keratinocytes and human and hamster melanoma cells (Slominski et al. 2002e) and enhanced the proliferation of epidermal keratinocytes and dermal fibroblasts (Bodo et al. 2010). Furthermore, TSH stimulated cyclic AMP production and altered the expression of several genes in human hair follicles and dermal papilla fibroblast cultures from normal female skin, however, did not affect hair growth and pigmentation (Bodo et al. 2009; van Beek et al. 2008). Expression of the TSH-R protein was detected in a wide panel of melanocytic lesions including melanoma. TSH activated the mitogen-activated protein kinase (MAPK) pathway and stimulated proliferation of melanoma cells, however, not melanocytes (Ellerhorst et al. 2010). Furthermore, TRH at low concentrations stimulated melanoma growth but not melanocytes' proliferation and its expression was increased in dysplastic nevi in contrast to benign nevi and was expressed in 63% of melanoma samples (Ellerhorst et al. 2006). Interestingly, other authors reported that suppression of MAP kinase and PI3K/Akt pathways, while leading to inhibition of cell proliferation, induced thyroid genes' expression including TSH-R and sodium/iodide symporter, which led to increased iodine uptake by melanoma cells (Hou et al. 2009).

Recent studies demonstrated the expression of thyroid factor-1, thyroglobulin (Bodo et al. 2009), and thyroperoxidase in the human skin (Cianfarani et al. 2010). T3 and T4 were also shown to modulate the expression of cytokeratins 6 and 14 genes



and downregulate TGF- $\beta$ 2 expression in hair follicles (van Beek et al. 2008). Thyroid hormones also stimulated hair pigmentation (van Beek et al. 2008) and mitochondrial function by upregulating the mRNA level of mitochondria-selective cytochrome-c-oxidase subunit 1 (MTCO1) and significantly increasing complex I and IV (cytochrome-c-oxidase) activities in the epidermis (Poeggeler et al. 2010).

It is well established that thyroid dysfunction alters skin physiology, but expression of the equivalent of HPT axis in the skin raises also the possibility of a cross-talk between local and systemic counterparts as it was demonstrated in amphibians (Vaudry et al. 1999). These interactions might have long-range consequences, especially for the regulation of global homeostasis, evolution of skin stress response systems, and development of thyroid-related autoimmune diseases (reviewed by Slominski et al. 2008b). From the clinical point of view, pathological exposure of TSH-R to immune cells in keratinocytes damaged by UVR or fibroblasts damaged during inflammation (Slominski et al. 2002e) can induce either production of anti-TSH-R antibodies leading to the uncontrolled stimulation of the thyroid gland or generation of anti-TSH-R clones of T lymphocytes—leading to immune destruction of the thyroid. These concepts, originally proposed by Slominski et al. (2002e), have been recently reemphasized defining a role of skin in thyroid autoimmune diseases (Cianfarani et al. 2010).

### 9.3 Conclusions

Different elements of the HPT axis are expressed in the skin and this expression is skin cell-type dependent. The expression of individual or networked HPT elements can regulate skin phenotype in a differentiated and context-dependent manner (Fig. 9.1). Possibly, communication between the cutaneous HPT axis and other local neuroendocrine networks as well as with central coordinating centers takes place and affects global homeostasis (Figs. 1.1 and 9.1).

# Chapter 10

## Cutaneous Opioid System

### 10.1 Overview

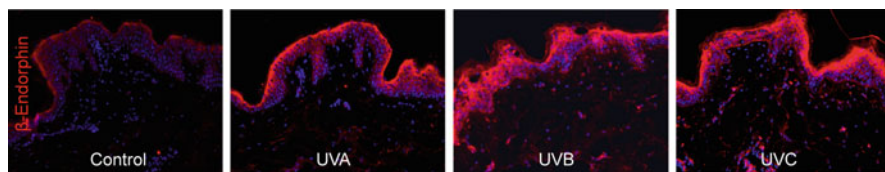
Endogenous opioid peptides derive from four different precursor proteins. Proopiomelanocortin (POMC) yields ACTH and endorphins, mainly  $\beta$ -endorphin ( $\beta$ -END). Proenkephalin (PENK) generates enkephalins (ENK), predominantly Met-enkephalin (MENK) and Leu-enkephalin (LENK). The proteolysis of prodynorphin (PDYN) results in the formation of dynorphin A (DYN A) and B (DYN B) (Przewlocki 2004; Przewlocki and Przewlocka 2005). Other POMC derivatives are endomorphins which are the cleavage products of a larger precursor molecule that yet has not been identified (Fichna et al. 2007).

Three classes of the Gi/Go/Gq-coupled opioid metabotropic receptor (OR) family have been identified: mu ( $\mu$ , MOR), delta ( $\delta$ , DOR), and kappa ( $\kappa$ , KOR). In addition, an orphan opioid-like nociceptin receptor (NOP), which has a 70% sequence homology with other opioid receptors, was also characterized (Jordan and Devi 1998; Jordan et al. 2000; Przewlocki and Przewlocka 2005; Salemi et al. 2005). Activation of ORs inhibits cAMP signaling as well as alters voltage-gated  $\text{Ca}^{2+}$  channel function and activates  $\text{K}^{+}$  inwardly rectifying channels. The endomorphins bind to MOR with the highest affinity, ENK preferentially binds to the DOR, and DYN favorably binds with KOR (Przewlocki 2004). The ligand-receptor affinity presents in the following order:  $\beta$ -END  $\mu$ ,  $\delta$  >  $\kappa$ ; LENK/MENK  $\delta$ ,  $\mu$ ; DYNA  $\kappa$  >  $\mu$  >  $\delta$  (Jordan et al. 2000; Tominaga et al. 2007).

### 10.2 Opioid Peptides in the Skin

#### 10.2.1 $\beta$ -Endorphin

It was shown that human and animal skin and/or cultured skin cells such as keratinocytes and melanocytes (normal and pathological) have the capability to transcribe and translate the precursor opioid protein genes (Nissen and Kragballe



**Fig. 10.1** Wavelength-dependent UV stimulation of  $\beta$ -END expression in the epidermal layer of human skin. CY<sup>TM</sup>3 positive (red) signals correspond to  $\beta$ -END immunoreactivity (methods described in Skobowiat et al. 2011)

1997; Polakiewicz et al. 1992; Salemi et al. 2005; Sikand et al. 2011; Skobowiat et al. 2011; Slominski et al. 1992, 1993b, 2000c, 2011c; Zagon et al. 1996). The POMC gene and protein and POMC-derived peptides (ACTH,  $\alpha$ -MSH, and  $\beta$ -END) were detected in epidermis, dermis, and adnexa and, additionally, can be released from the cutaneous nerve endings (Hasse et al. 2007; Mazurkiewicz et al. 2000; Slominski et al. 1992, 1993b, 1998a, 2000c; Tobin and Kauser 2005a; Wintzen and Gilchrest 1996). The cutaneous expression of POMC was first found in rodent (hamster and mouse) melanomas (Slominski 1991) and in mouse (Slominski et al. 1992) and human (Slominski et al. 1993c) skin, where the  $\beta$ -END antigen was detected as well.  $\beta$ -endorphin stimulates keratinocyte migration in vitro (Tominaga et al. 2007), induces epidermal and follicular melanogenesis (Kauser et al. 2004), and also controls hair growth, wound healing, and cellular differentiation (Schmelz and Paus 2007). Furthermore,  $\beta$ -END increases and modulates the number of dendritic processes of hair follicle melanocytes (Kauser et al. 2004). Some authors reported that plasma  $\beta$ -END levels increase after UV exposure, which would explain the euphoric behavior observed in beachgoers (Fallahzadeh and Namazi 2009). Other authors could not observe this correlation (Wintzen et al. 2001). Recently, UVR-induced increase of  $\beta$ -END level has been observed in ex vivo maintained human skin (Skobowiat et al. 2011) (Fig. 10.1).

### 10.2.2 *Proenkephalin*

The expression of the *PENK* gene and the subsequent processing of PENK to MENK/LENK (in a cell type-dependent manner) were demonstrated in the skin by RT-PCR, Western blotting, immunocytochemistry, time-of-flight/liquid chromatography, and mass spectroscopy (Slominski et al. 2011c). PENK immunoreactivity was markedly restricted to differentiating keratinocytes of the stratum spinosum and granulosum, whereas proliferating basal keratinocytes did not exhibit this immunoreactivity (Slominski et al. 2011c; Zagon et al. 1996). Furthermore, physical (UVB) and biological (lipopolysaccharide) stressors demonstrated cell type-specific time- and dose-dependent stimulation of *PENK* gene expression (Slominski et al. 2011c). Also, fetal mesenchymal skin cells expressed and produced significant amount of PENK, indicating its association with cell proliferation

(Polakiewicz et al. 1992). MENK released from rat Merkel cells acted in an autocrine/paracrine fashion by the inhibition of cell granules' release via decrease of intracellular  $\text{Ca}^{2+}$  concentration (Tachibana and Nawa 2005). ENK plays a role in the differentiation of epidermal keratinocytes and has direct antimicrobial activities which contribute to the skin protective barrier against noxious factors (Nissen and Kragballe 1997; Slominski et al. 2011c).

### ***10.2.3 Dynorphins***

PDYN and DYN A are expressed in human skin cells as well as in cutaneous nerve fibers (Grando et al. 1995; Hassan et al. 1992; Salemi et al. 2005; Taneda et al. 2011; Tominaga et al. 2007). They are responsible for pilomotor activity (Gibbins 1992) and nociception. Enhanced production of DYN A was found in atopic dermatitis (Tominaga et al. 2007). Endomorphin 1 and 2 immunoreactivity was found in nerve fibers of the rat skin; however, the physiological function of these peptides still has to be determined (Barr and Zadina 1999; Borzsei et al. 2008).

## **10.3 Opioid Receptors and Their Function in the Skin**

Previous studies showed that not only opioid peptides but also their receptors were expressed in the skin (Grando et al. 1995; Nissen and Kragballe 1997; Salemi et al. 2005; Tachibana and Nawa 2005). These findings define skin as an active environment for opioid action. Ectoderm-derived cells, e.g., keratinocytes and melanocytes, express ORs, however, at much lower level (by a factor of 200–20,000) than neurons. MOR showed a stronger expression than DOR in keratinocytes, but an opposite expression pattern was found in mesenchyme-derived fibroblasts (Bigliardi et al. 1998). Furthermore, KOR was detected in fibroblasts and mononuclear blood cells of normal human skin and DOR was expressed in fibroblasts isolated from human skin (Salemi et al. 2005). Enhanced expression of DOR and KOR in the skin justifies the exploration of novel  $\delta$  and  $\kappa$  acting compounds as specific targets for future opioid therapy. OR-induced signaling can affect cell differentiation, migration, as well as cytokeratin and cytokine expression in human epidermis. Thus, opioid receptors may be involved not only in the regulation of normal skin homeostasis but also in wound healing and scar formation (Bigliardi et al. 2009).

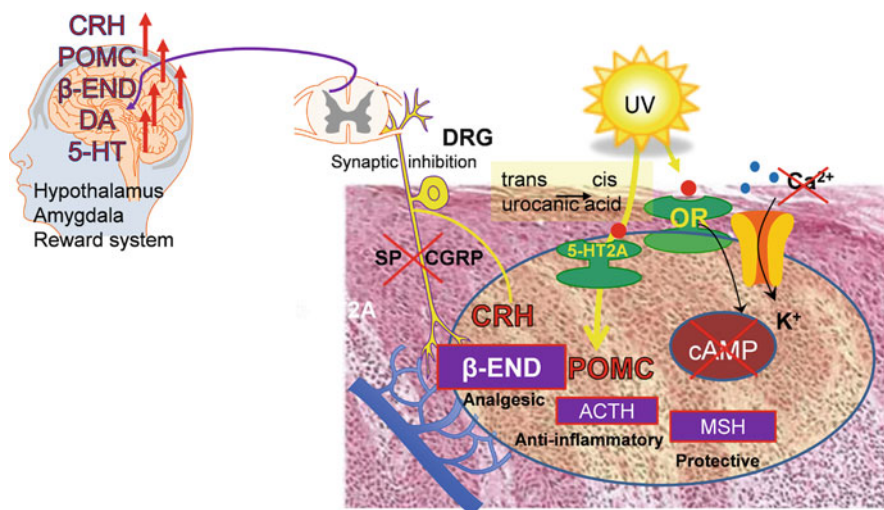
Opioids are best known for their antinociception, which can also be initiated by the activation of ORs outside the central nervous system (Salemi et al. 2005). Their inhibitory activities are related to the ligand activation of  $\text{Gi}, \text{o}, \text{q}$  proteins which downregulates adenylyl cyclase activity and, eventually, inhibits protein phosphorylation (Chizhnikov et al. 2005). In addition, modifications of ion channels' activity have further expanded the spectrum of opioid-induced biological action. All ORs

couple to various  $\text{Ca}^{2+}$  channels and are known to inhibit their activity. Stimulation of ORs also increases potassium conductance across the cellular membranes (Jordan and Devi 1998; Jordan et al. 2000). Opiates increase the release of dopamine via presynaptic inhibition of GABA release, which leads to a reward response, an effect observed in tanning addicts (Harrington et al. 2006; Nolan and Feldman 2009; Przewlocki 2004). Inhibitory action of opiates on noradrenergic activity is mediated by MOR, and its activation enhances CRF production and release (Armario 2010). Opioids exert immunomodulatory effects in peripheral tissues including skin by stimulating lymphocyte proliferation, antibody production, T and NK cell activity, and chemotaxis of macrophages and granulocytes (Fallahzadeh and Namazi 2009; Jankowska and Schomburg 1998; Jordan and Devi 1998; Mousa et al. 2007).

Opioids produced by skin cells may probably act as para- and autocrine modulators that influence gene expression. Additionally, they may enter the systemic circulation being transported by dermal veins, and interact with specific receptors localized on cutaneous nerve fibers (Bigliardi et al. 2009; Borzsei et al. 2008; Slominski and Wortsman 2000; Slominski et al. 2000c). The last property is required for their analgesic and anti-inflammatory activity. Transduced nervous signals from peripheral tissues could be conveyed to dorsal root ganglia (DRG) sensory neurons. Thus, via synaptic inhibition neuropeptides like substance P (SP) and calcitonin gene-related peptide (CGRP), responsible for pain and inflammation, would be downregulated and not delivered to skin (Borzsei et al. 2008; Slominski and Wortsman 2000; Tobin 2006). Furthermore, the ascending activator signals could be transferred via the nucleus of the solitary tract (NTS) to thalamus, hypothalamus, especially to paraventricular and arcuate nuclei, and amygdala where they could exert their systemic actions (Slominski et al. 2008b) and enhance the activity of the reward system (Armario 2010; Przewlocki 2004; Przewlocki and Przewlocka 2005; Slominski 2005; Slominski and Wortsman 2000) (Fig. 10.2).

## 10.4 Opioid System and Pruritus

It has been known for decades that analgesia resulting from MOR activation induces itch, whereas MOR antagonists, such as naltrexone, inhibit itch (Yosipovitch 2010). It is widely accepted that the activation of KOR signaling suppresses, while that of MOR stimulates, itch (Bigliardi et al. 2009; Roosterman et al. 2006). This has led to the opioidergic system being targeted by new antipruritic medications (Patel and Yosipovitch 2010; Schmelz 2010). A novel KOR agonist, nalfurafine/TRK-820, revealed antipruritic activity in morphine-, histamine-, and substance P-induced animal scratching models (Ko and Husbands 2009). The number of nerve fibers entering the epidermis tended to increase in approximately 40% of psoriatic patients claiming itch sensation compared to healthy controls (Taneda et al. 2011). There were no differences in epidermal number of MOR and  $\beta$ -END amount; however, levels of KOR and DYN A were



**Fig. 10.2** Hypothetical pathway of addictive activities of the cutaneous neuroendocrine system induced by UV stimulation

significantly decreased between healthy controls and psoriatic patients (Taneda et al. 2011). Interestingly, opioids were shown to act upon capsaicin-sensitive nerve fibers and inhibit the release of inflammatory neuropeptides such as SP, neurokinin A, and CGRP, i.e., neuropeptides that are indirectly involved in eliciting pruritus by releasing the pruritogen—histamine (Roosterman et al. 2006; Stander et al. 2002).

It was shown that UV-induced keratinocyte-delivered nerve growth factor (NGF), upon retrograde transport from skin toward dorsal root ganglion, enhanced the expression of neuropeptides SP and CGRP and upregulated the number of MORs localized on cutaneous sensory nerve fibers (Mousa et al. 2007; Roosterman et al. 2006). Recent studies have provided evidence that, indeed, there are itch-specific receptors in the skin. Using in vitro binding assays, it was observed that the proteolytically cleaved product of proenkephalin A, the bovine adrenal medulla peptide 8–22 (BAM8–22), potently activated the Mas-related G-protein-coupled receptors (Mrgprs) in an opioid-independent mechanism (Sikand et al. 2011).

## 10.5 Opioids and Addiction

UV exposure during indoor tanning damages DNA, thereby leading to premature skin aging and the development of skin cancer as well as malignant melanoma (Harrington et al. 2006). The majority of beachgoers reported behaviors consistent with those of an addictive disorder like continuation of tanning despite attempts to stop, persistent tanning in the presence of adverse reactions, and the neglect of other

responsibilities in order to maintain a tan (Keen et al. 2008). Psychological dependence is suggested by tanners' reports of relaxation and positive mood effects as a result of UV exposure. These observations may be explained by UV-induced production of cutaneous  $\beta$ -END (Skobowiat et al. 2011) resulting from local transcription, translation, and further cleavage of POMC leading to the production of  $\beta$ -END (reviewed by Slominski 2003; Slominski et al. 1993b, 2000c).

Solar UV energy adsorbed by the epidermis also results in the transformation of a chromophore-like trans-urocanic acid to its cis-isomer, which reportedly could have an agonistic activity on serotonin receptor 2A (5HT2A) (Walterscheid et al. 2006). Thus, UV light can also alter cutaneous serotonergic system with subsequent effects on the CNS affecting the mood (Harrington et al. 2011; Kourosh et al. 2010; Slominski et al. 2005c). Upon UV exposure, 5-HT could be transported from the skin via its blood vessels or activate ascending nerve fibers which affect brain activity (Nordlind et al. 2008; Slominski et al. 2005c). Specifically, brain striatal regions, including the nuclei accumbens, caudate and putamen, could be activated during UVR exposure (reviewed by Kourosh et al. 2010). Frequent tanning may involve CNS reward and/or reinforcement over the often-stated goal of "getting a tan." The ventral striatum (or nucleus accumbens) activation is typically associated with drug-induced reward observed in cocaine and nicotine smokers (Harrington et al. 2006). In fact, the term "psychodermatology" is being used to describe the mind-skin connection (Reich et al. 2010). From an evolutionary perspective, it is reasonable to suggest that sunlight may have central rewarding properties (Figs. 1.1 and 10.2) given the importance for human health of UV-mediated vitamin D synthesis.

## 10.6 Conclusions

Opioids constitute a heterogeneous family of active peptides which play important roles in cutaneous nociception, immunomodulation, signal transduction, and evoking or attenuating of pruritus, depending on differential receptor activation. Their main inhibitory properties are related to the inhibition of cell membrane calcium channels that suppress the release of proinflammatory and pain transmitters like SP and CGRP. Both exogenously applied and endogenous opioids interact with a whole range of receptors and contribute to the neuroendocrinological functions of the skin, also at the systemic level (Figs. 1.1, 1.2 and 10.2).

# Chapter 11

## Cutaneous Endocannabinoid System

### 11.1 Overview

Endocannabinoids (ECS) constitute lipid mediators (amides, esters, and ethers of long chain polyunsaturated fatty acids) which act similarly to the exogenous  $\Delta^9$  tetrahydrocannabinol (THC; the main psychoactive ingredient of the plant *Cannabis sativa*) and are produced in humans and animals (Maccarrone et al. 2003; Rahn and Hohmann 2009). The cutaneous ECS system is fully functional due to the expression of ECS and their receptors as well as ECS-degrading enzymes. All the components of the skin ECS system were shown to modulate the proliferation, differentiation, growth, and apoptosis of various skin cell types as well as tumorigenesis and local cytokine production (reviewed by Biro et al. 2009; Kupczyk et al. 2009; Toth et al. 2011). ECS are synthesized “on demand” by receptor-stimulated cleavage of membrane lipid precursors and are not stored in synaptic vesicles which distinguishes them from typical neurotransmitters. ECS reuptake may be facilitated by a transporter that has not been cloned yet; however, pharmacological inhibitors of ECS transport have nonetheless been developed (Guindon and Hohmann 2009). The lipophilic nature of ECS allows them to activate various enzymes in cytosol and membraneous compartments (Kupczyk et al. 2009).

The most extensively studied ECS are *N*-arachidonylethanolamide (anandamide/AEA), 2-arachidonoylglycerol (2-AG), and *N*-palmitoylethanolamide (PEA), while their main catabolic enzymes are fatty acid amid hydrolase (FAAH) and the monoacylglycerol lipase (MAGL).



## 11.2 Endocannabinoid Receptors and Mechanisms of ECS Action

Two receptor types for cannabinoids, CB1 and CB2, have been identified beyond doubt; however, some researchers assume the existence of a third one, CB3, which has not been cloned yet (Kupczyk et al. 2009). The amino acid sequences of the CB receptors are conserved in many species, including humans (Kupczyk et al. 2009). CB1 is predominantly expressed in the central nervous system and other tissues, while CB2 was mainly found in non-neuronal cells and tissues related to the immune system, like lymphocytes, macrophages, spleen, and thymus. In the skin, both receptor types are expressed on nerve endings, mast cells, keratinocytes, and adnexa (Biro et al. 2009; Maccarrone et al. 2003). Classical activation of CBs, which belong to the coupled Gi/o family of G proteins, inhibits N- and P/Q-type  $\text{Ca}^{2+}$  channels, activates A-type, inwardly rectifying potassium conductance channels, and inhibits M-type potassium channels. Furthermore, binding of CB2 receptor ligands modulates MAPK activity resulting ultimately in decreased cAMP production and suppression of neuronal excitability and transmitter release (reviewed by Irving et al. 2002). Additionally, CB1 activation can inhibit conductance of serotonin receptor 3 (5-HT3) ion channels, modulate the production of NO, alter voltage-sensitive sodium channel activity, and activate the  $\text{Na}^+/\text{H}^+$  exchanger (Guindon and Hohmann 2009; Rahn and Hohmann 2009). It was shown that CB1 and transient receptor potential vanilloid-1 (TRPV-1) are co-localized on sensory nerve endings in the skin (Akerman et al. 2004) which suggests that ECS can also activate TRPV-1 receptors in the skin (Karst et al. 2010; Maccarrone et al. 2003). Anandamide acts at vanilloid receptors and blocks directly the background  $\text{K}^+$  channel affecting NMDA transmission in the brain (reviewed by Irving et al. 2002). CB1 receptors are localized presynaptically on GABA-ergic and glutamatergic interneurons (Kupczyk et al. 2009). CB1 activation results in a decreased release of neurotransmitters such as GABA ( $\gamma$ -aminobutyric acid) and glutamate. This retrograde signaling mechanism suggests an important modulatory role of ECS in controlling neuronal excitability and homeostasis. ECS mediate habituation to stress by restraining HPA axis response and maintaining body's homeostasis (Finn 2010).

## 11.3 Effects of ECS on Proliferation and Differentiation of Keratinocytes

CB1 agonists inhibited proliferation of cultured human epidermal keratinocytes and anandamide (AEA) markedly suppressed cell growth and induced dose- and CB1-dependent apoptosis in human HaCaT keratinocytes (Paradisi et al. 2008; Toth et al. 2011). AEA inhibited hair shaft elongation and proliferation of hair matrix keratinocytes (Telek et al. 2007). Cannabinoids also induced intraepithelial

apoptosis and premature hair follicle regression (characteristic signs of catagen transformation in hair follicles)—processes that were inhibited by a selective CB1 antagonist (Pucci et al. 2011; Toth et al. 2011). CB1 is expressed in a hair cycle-dependent manner and negatively regulates human hair growth in an autocrine—paracrine manner. Indeed, it was shown that CB1 antagonists induced hair growth in mice (Srivastava et al. 2009).

Differentiating human keratinocytes had decreased levels of endogenous AEA due to increased degradation of this lipid (Maccarrone et al. 2003). Moreover, it was shown that exogenous AEA inhibited keratinocyte differentiation *in vitro* acting via a CB1-dependent mechanism that involved inactivation of protein kinase C, activating protein-1 (AP-1), and transglutaminase (Maccarrone et al. 2003). High expression of CB1 in epidermal granular and spinous layers suggests participation of ECS in keratinocytes' differentiation (Stander et al. 2005).

## 11.4 Effects on Tumorigenesis

Various human skin tumors (e.g., basal and/or squamous cell carcinoma) express both CB1 and CB2 receptors (Casanova et al. 2003; Zheng et al. 2008). Local administration of synthetic CB1 and CB2 agonists inhibited growth of skin malignant tumors in nude mice by increasing intra-tumor apoptosis and impairing tumor vascularization (Casanova et al. 2003). ECS also inhibited *in vivo* growth of mouse melanomas that expressed CB1 and CB2 by decreasing growth, proliferation, angiogenesis, and metastasis formation, while increasing apoptosis (Blazquez et al. 2006). It was also found that human squamous cell carcinoma overexpressed CB2 at the both mRNA and protein levels (Zhao et al. 2010). UVB exposure of experimentally induced papilloma in mouse skin led to a local activation of CB1/2 receptors. While the absence of the functional CB1/2 receptors in double knock-out mice resulted in a resistance to UVB-induced inflammation and a marked decrease in UVB-induced skin carcinogenesis. Thus, the CB1/2 receptors play a key role in UV-induced inflammation and skin cancer development (Zheng et al. 2008).

## 11.5 Effects on Inflammation

CB2 receptor activation, in general, mediates immunosuppressive effects, which limit inflammation and associated tissue injury in large number of pathological conditions. Interestingly, stimulation of CB2 receptors in immune cells after initial decrease in cAMP production may lead to a sustained, pronounced increase in cAMP levels, which results in the suppression of T-cell receptor signaling through a cAMP/PKA/Csk/Lck pathway (reviewed by Pacher and Mechoulam 2011). Recent studies have also revealed that ECS may affect proliferation and apoptosis of T and

B lymphocytes, inflammatory cytokine production, and immune cell activation by inflammatory stimuli (e.g., LPS), macrophage-mediated killing of sensitized cells, chemotaxis, and inflammatory cell migration (Sanchez and Garcia-Merino 2012). In a mouse model of contact allergy, the cutaneous inflammation has been suppressed by local administration of  $\Delta^9$  tetrahydrocannabinol (THC) and CB agonists (Karsak et al. 2007). In murine dermatitis elevated 2-AG (2-arachidonoylglycerol) levels were observed, and inflammatory symptoms were markedly attenuated by CB2 (but not CB1) agonists (Oka et al. 2006). Decreased dermal fibrosis (bleomycin-induced) and inflammation were observed upon treatment with a CB2 agonist, suggesting a potential therapeutic application of selective CB2 agonists in early inflammatory stages at the local and systemic levels (Akhmetshina et al. 2009).

## 11.6 Effects on Pain Sensation

Antinociceptive activity of cannabinoids has been one of the main reasons of their worldwide usage in many communities. Recently described mechanisms of antinociceptive action of ECS involve the specific CB's activation as well as interaction with other receptors and pathways related to pain sensation at the central (Connell et al. 2006; Irving et al. 2002) and peripheral (Finn 2010; Karst et al. 2010; Amaya et al. 2006) levels. Activation of CBs changes cellular  $\text{Ca}^{2+}$  and  $\text{K}^{+}$  conductance and decreases cAMP levels leading to an inhibition of neurotransmitter's release which is involved in pain sensation (Rahn and Hohmann 2009; Akerman et al. 2004).

Antinociceptive effects of CB1/2 agonists, and FAAH inhibitors, which prolong the action of ECS, have been reported. It was shown in rats that CB2 agonists stimulate release of  $\beta$ -END from keratinocytes, at the local mu-opioid related fashion, to inhibit nociception in the skin (Ibrahim et al. 2005). The application of electroacupuncture and CB2 agonist reduced inflammatory pain due to increased POMC mRNA and  $\beta$ -END levels (via mu-opioid receptor mechanism) in inflamed rat skin (Su et al. 2011). Increased mobilization and activity of ECS in amygdala, observed after foot shock stress in rats, lead to an attenuation of nociceptive pathways due to CB1 activation (Connell et al. 2006). Furthermore, cannabis (inhaled as aerosol) augmented opioid analgesia through a pharmacodynamic mechanism, in experimental study performed in humans (Abrams et al. 2011). Elevated levels of ECS in models of chronic pain are likely to counteract increased neuronal activity driven by afferent nociceptive inputs. ECS-induced inhibition in neurotransmission may modulate central sensitization observed during pain stimuli (Connell et al. 2006; Abrams et al. 2011, reviewed by Karst et al. 2010). The cutaneous application of synthetic CB agonist was shown to reverse inflammatory thermal hyperalgesia evoked by Freund's adjuvant application in rats (Amaya et al. 2006). Elevated ECS level, evoked by blocking of catabolic (mainly FAAH) enzymes activity, is effective and desired feature in decreasing pain perception

during osteoarthritis in rats (Sagar et al. 2010). Extended ECS action, due to FAAH deactivation, is likely to be more beneficial compared to direct activation of CB1 receptors (reviewed by Karst et al. 2010; Rahn and Hohmann 2009). Interestingly, both AEA and 2-AG can be metabolized by cyclooxygenase-2, which may contribute to the pain relieving properties of nonsteroidal anti-inflammatory drugs that act by inhibiting the cyclooxygenase (Rahn and Hohmann 2009).

## 11.7 Antipruritic Effects

Pruritus, an unpleasant cutaneous sensation associated with an immediate desire to scratch, may be interpreted as one of the body's defense mechanisms (reviewed by Steinhoff et al. 2006). Pruriceptor stimulation conveys the transduced signal via histamine-positive mechano-insensitive C-fibers originating in DRG up to itch-selective units in lamina I of the spinal cord. Ascending signals project via posterior part of the ventromedial thalamic nucleus to finally reach the dorsal insular cortex (reviewed by Steinhoff et al. 2006). Inflammatory mediators released after disruption of cutaneous barrier function or UV radiation can stimulate sensory nerve endings (reviewed by Roosterman et al. 2006; Slominski and Wortsman 2000), and thus, induce pruritus (reviewed by Yosipovitch 2010). Peripheral administration of cannabinoid receptor agonists attenuated histamine-induced itch in humans (Dvorak et al. 2003). Recent study with the use of *N*-palmitoylethanolamine (cannabinoid-like amide acting through PPAR- $\alpha$ ), which was added as a component of itch-relieving creams, alleviated pruritus in patients with atopic dermatitis, lichen simplex, and prurigo nodularis (Kircik 2010). These promising preliminary results suggest that new therapies targeting cannabinoid receptors may result in providing effective antipruritic medication in the future.

## 11.8 Conclusions

The recently described endocannabinoid system contributes to the abundant neuro-endocrine activities of the skin. ECS participate in a number of pathophysiological processes in the skin and present there profound anti-inflammatory, anti-tumorigenic, antinociceptive, and antipruritic effects. ECS interact with two specific receptors which lead to synaptic inhibition of many neurotransmitter systems. Furthermore, ECS can interact with other receptors (opioid, serotonin, and TRPV) by nonspecific binding, modulating the release of other neurotransmitters and hormones. Thus, cutaneous ECS can participate in the regulation of local and systemic homeostasis (Figs. 1.1 and 1.2).

## Chapter 12

# Perspectives

Described as the body's largest organ, the skin is strategically located at the interface with the external environment where it has evolved to detect, integrate, and respond to a diverse range of stressors including UV radiation. Recent findings have established the skin as a peripheral neuroendocrine organ that is tightly networked to central stress axes (Fig. 1.2). This capability contributes to the maintenance of skin's and body's homeostasis. Specifically, epidermal and dermal cells produce and respond to classical stress neurotransmitters, neuropeptides, and hormones, and this production is modified by ultraviolet radiation and biological, chemical, and physical factors. Examples of potent epidermal products include biogenic amines (catecholamines, serotonin, and *N*-acetyl-serotonin) (Figs. 2.1–2.3), acetylcholine, melatonin and its metabolites (Figs. 2.5 and 3.1), proopiomelanocortin-derived ACTH,  $\beta$ -endorphin and MSH peptides, corticotropin-releasing factor and related urocortins (Figs. 5.1, 5.2, 7.3 and 7.4), corticosteroids and their precursor molecules, thyroid-related hormones (Fig. 9.1), opioids, and cannabinoids. The production of these molecules in the skin is hierarchical, following the algorithms of classical neuroendocrine axes (e.g., hypothalamic pituitary adrenal axis (HPA), hypothalamic–thyroid axis, serotonergic/melatonergic, catecholaminergic and cholinergic systems). The deregulation of these systems may be involved in the etiology of some skin diseases. These local neuroendocrine systems represent exquisite regulatory levels addressed at restricting the effect of noxious agents to preserve local and, consequently, global body's homeostasis and adapt to changing external environment. Furthermore, the skin-derived signals may also activate cutaneous sensory nerve endings to alert the brain on environment- or pathology-induced changes in the epidermal and dermal milieu, or alternatively, to activate other coordinating centers by spinal cord neurotransmission with or without brain's involvement (Fig. 1.1). Finally, the local neuroendocrine system will imprint resident and circulating immune cells to act as cellular messengers sent to other organs to coordinate responses to the changing environment (1.1).

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